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OM protein - protein search, using sw model

Run on: January 28, 2006, 06:20:07 ; Search time 18.6667 Seconds
(without alignments)
53.149 Million cell updates/sec

Title: US-09-897-042-21

Perfect score: 68

Sequence: 1 KISFPEGPPKY 12

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents AA:*

- 1: /cgn2_6/prodata/1/1aa/5 COMB.pep.*
- 2: /cgn2_6/prodata/1/1aa/6 COMB.pep.*
- 3: /cgn2_6/prodata/1/1aa/H COMB.pep.*
- 4: /cgn2_6/prodata/1/1aa/ECTUS COMB.pep.*
- 5: /cgn2_6/prodata/1/1aa/RE COMB.pep.*
- 6: /cgn2_6/prodata/1/1aa/backfiles1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	ID	Description
1	68	100.0	160	1	US-07-847-010-23
2	57	83.8	160	1	US-07-847-010-3
3	42	61.8	225	2	US-09-866-538-12
4	42	61.8	225	2	US-09-865-291-12
5	42	61.8	240	2	US-10-152-296-2
6	41	60.3	98	2	US-09-205-658-194
7	41	60.3	109	2	US-09-513-999C-6109
8	41	60.3	122	2	US-09-205-658-175
9	41	60.3	152	2	US-09-216-430C-20
10	41	60.3	153	1	US-08-748-703-1
11	41	60.3	153	2	US-09-132-861-1
12	41	60.3	153	2	US-09-949-016-6465
13	41	60.3	168	2	US-09-949-016-10584
14	41	60.3	621	2	US-09-331-568A-2
15	41	60.3	621	2	US-09-331-568A-26
16	41	60.3	632	2	US-09-205-658-159
17	41	60.3	636	2	US-09-205-658-160
18	40	58.8	259	1	US-08-861-269-1
19	40	58.8	259	1	US-09-134-596-1
20	40	58.8	259	2	US-09-293-273-1
21	40	58.8	284	2	US-09-270-767-46828
22	39	57.4	55	2	US-09-621-976-7597
23	39	57.4	514	2	US-08-688-988-35
24	38	55.9	112	2	US-09-583-110-3506
25	38	55.9	350	2	US-09-248-796A-19957
26	38	55.9	435	2	US-09-107-433-2745
27	38	55.9	524	2	US-08-688-988-34

28	38	55.9	525	2	US-08-688-988-10	Sequence 10, Appl
29	38	55.9	1049	2	US-09-107-532A-5966	Sequence 5966, Ap
30	38	55.9	1751	2	US-09-136-574A-44	Sequence 44, Appl
31	37.5	55.1	352	2	US-09-252-991A-20145	Sequence 20145, A
32	37	54.4	15	1	US-08-080-073-4	Sequence 4, Appl
33	37	54.4	15	1	US-08-080-073-18	Sequence 18, Appl
34	37	54.4	15	1	US-08-080-073-22	Sequence 22, Appl
35	37	54.4	40	2	US-09-277-716-25	Sequence 25, Appl
36	37	54.4	40	2	US-09-609-161B-25	Sequence 25, Appl
37	37	54.4	76	2	US-09-293-505-4	Sequence 4, Appl
38	37	54.4	76	2	US-09-060-939A-4	Sequence 4, Appl
39	37	54.4	122	2	US-09-513-999C-7924	Sequence 7924, Ap
40	37	54.4	137	2	US-09-513-999C-4903	Sequence 4903, Ap
41	37	54.4	154	1	US-08-464-342-4	Sequence 4, Appl
42	37	54.4	154	1	US-08-748-703-3	Sequence 3, Appl
43	37	54.4	154	1	US-08-464-604A-4	Sequence 4, Appl
44	37	54.4	154	1	US-08-875-272-4	Sequence 4, Appl
45	37	54.4	154	1	US-08-718-538-2	Sequence 2, Appl

ALIGNMENTS

RESULT 1
US-07-847-010-23
; Sequence 23, Application US/07847010
; Patent No. 5693495
; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikertorfer, Arnold
; APPLICANT: Valenta, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
; APPLICANT: Rumpold, Helmut
; APPLICANT: Scheiner, Otto
; APPLICANT: Ebner, Christof
; APPLICANT: Ferreira, Fatima
; TITLE OF INVENTION: Allergens of Alder Pollen and
; TITLE OF INVENTION: Applications Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 23:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:

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; ORGANISM: birch (Betula sp.)
; IMMEDIATE SOURCE:
; LIBRARY: POLLEN FROM ALLERCON AB, ENGELHOLM, SWEDEN
US-07-847-010-23

Query Match          100.0%; Score 68; DB 1; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.00016;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGPFKY 12
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Db 56 KISFPEGPFKY 67

RESULT 2
US-07-847-010-3
; Sequence 3, Application US/07847010
; Patent No. 5693495
; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikerstorfer, Arnold
; APPLICANT: Valenta, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
; APPLICANT: Rumpold, Helmut
; APPLICANT: Scheiner, Otto
; APPLICANT: Ebner, Christof
; APPLICANT: Ferreira, Fatima
; TITLE OF INVENTION: Allergens of Alder Pollen and
; TITLE OF INVENTION: Applications Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; ORGANISM: Alder (Alnus sp.)
US-07-847-010-3

Query Match          83.8%; Score 57; DB 1; Length 160;
Best Local Similarity 83.3%; Pred. No. 0.015;
Matches 10; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KISFPEGPFKY 12
    |||||
Db 56 KISFPEGPFKY 67

; ORGANISM: birch (Betula sp.)
; IMMEDIATE SOURCE:
; LIBRARY: POLLEN FROM ALLERCON AB, ENGELHOLM, SWEDEN
US-07-847-010-23

Query Match          100.0%; Score 68; DB 1; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.00016;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGPFKY 12
    |||||
Db 56 KISFPEGPFKY 67

RESULT 3
US-09-866-538-12
; Sequence 12, Application US/09866538
; Patent No. 6852849
; GENERAL INFORMATION:
; APPLICANT: REGENTS OF THE UNIVERSITY OF CALIFORNIA
; APPLICANT: TSJEN, Roger
; APPLICANT: Campbell, Robert
; TITLE OF INVENTION: NON-OLIGOMERIZING FLUORESCENT PROTEINS
; FILE REFERENCE: REGEN1530-2
; CURRENT APPLICATION NUMBER: US/09/866,538
; CURRENT FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 29
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 12
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Discosoma sp.
US-09-866-538-12

Query Match          61.8%; Score 42; DB 2; Length 225;
Best Local Similarity 63.6%; Pred. No. 9.8;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KISFPEGPFKY 11
    |||||
Db 84 KLSFPEGFKWE 94

RESULT 4
US-09-865-291-12
; Sequence 12, Application US/09865291
; Patent No. 6900304
; GENERAL INFORMATION:
; APPLICANT: REGENTS OF THE UNIVERSITY OF CALIFORNIA
; APPLICANT: TSJEN, Roger
; APPLICANT: TING, Alice
; APPLICANT: ZHANG, Jin
; TITLE OF INVENTION: EMISSION RATIONOMETRIC INDICATORS OF PHOSPHORYLATION
; FILE REFERENCE: REGEN1550
; CURRENT APPLICATION NUMBER: US/09/865,291
; CURRENT FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 42
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 12
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Discosoma sp.
US-09-865-291-12

Query Match          61.8%; Score 42; DB 2; Length 225;
Best Local Similarity 63.6%; Pred. No. 9.8;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KISFPEGPFKY 11
    |||||
Db 84 KLSFPEGFKWE 94

RESULT 5
US-10-152-296-2
; Sequence 2, Application US/10152296
; Patent No. 6723537
; GENERAL INFORMATION:
; APPLICANT: Peelle, Beau
; APPLICANT: Rigel Pharmaceuticals, Incorporated
; TITLE OF INVENTION: Directed Evolution of Protein in Mammalian Cells
; FILE REFERENCE: 021044-000110US
; CURRENT APPLICATION NUMBER: US/10/152,296
; CURRENT FILING DATE: 2002-12-10
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; PRIOR APPLICATION NUMBER: US 60/291,871
; PRIOR FILING DATE: 2001-05-18
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 2
; LENGTH: 240
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:mammalian
; OTHER INFORMATION: codon-optimized variant (DsRED) of Discosoma sp.
; OTHER INFORMATION: "red" red fluorescent protein (RFP)
US-10-152-296-2

Query Match 61.8%; Score 42; DB 2; Length 240;
Best Local Similarity 63.6%; Pred. No. 11;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KISFPEGPPFK 11
|:|||||: :
Db 85 KLSFPEGPKWE 95

RESULT 6
US-09-205-658-194
; Sequence 194, Application US/09205658
; Patent No. 6861256
; GENERAL INFORMATION:
; APPLICANT: Ruvkun, Gary
; TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
; TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
; FILE REFERENCE: 00786/351004
; CURRENT APPLICATION NUMBER: US/09/205,658
; CURRENT FILING DATE: 1998-12-03
; EARLIER APPLICATION NUMBER: 08/857,076
; EARLIER FILING DATE: 1997-05-15
; EARLIER APPLICATION NUMBER: 08/888,534
; EARLIER FILING DATE: 1997-07-07
; EARLIER APPLICATION NUMBER: US98/10080
; EARLIER FILING DATE: 1998-05-15
; NUMBER OF SEQ ID NOS: 328
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 194
; LENGTH: 98
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-205-658-194

Query Match 60.3%; Score 41; DB 2; Length 98;
Best Local Similarity 100.0%; Pred. No. 5.8;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 SPPEGFP 9
|:|||||
Db 68 SPPEGFP 74

RESULT 7
US-09-513-999C-6109
; Sequence 6109, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Duclert, A.
; APPLICANT: Giordano, J.Y.
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; FILE REFERENCE: 59.US2.REG
; CURRENT APPLICATION NUMBER: US/09/513,999C
; CURRENT FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/122,487
; PRIOR FILING DATE: 1999-02-26

; NUMBER OF SEQ ID NOS: 36681
; SOFTWARE: Patent.pm
; SEQ ID NO 6109
; LENGTH: 109
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: 105
; OTHER INFORMATION: Xaa=Leu or Pro
US-09-513-999C-6109

Query Match 60.3%; Score 41; DB 2; Length 109;
Best Local Similarity 63.6%; Pred. No. 6.5;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 KISFPEGPPFK 11
|:|||||: :
Db 54 RISFPPEYFPK 64

RESULT 8
US-09-205-658-175
; Sequence 175, Application US/09205658
; Patent No. 6861256
; GENERAL INFORMATION:
; APPLICANT: Ruvkun, Gary
; APPLICANT: Ogg, Scott
; TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
; TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
; FILE REFERENCE: 00786/351004
; CURRENT APPLICATION NUMBER: US/09/205,658
; CURRENT FILING DATE: 1998-12-03
; EARLIER APPLICATION NUMBER: 08/857,076
; EARLIER FILING DATE: 1997-05-15
; EARLIER APPLICATION NUMBER: 08/888,534
; EARLIER FILING DATE: 1997-07-07
; EARLIER APPLICATION NUMBER: US98/10080
; EARLIER FILING DATE: 1998-05-15
; NUMBER OF SEQ ID NOS: 328
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 175
; LENGTH: 122
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-205-658-175

Query Match 60.3%; Score 41; DB 2; Length 122;
Best Local Similarity 100.0%; Pred. No. 7.4;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 SPPEGFP 9
|:|||||
Db 38 SPPEGFP 44

RESULT 9
US-09-216-430C-20
; Sequence 20, Application US/09216430C
; Patent No. 6734283
; GENERAL INFORMATION:
; APPLICANT: Chau, Vincent
; TITLE OF INVENTION: Human Proteins Responsible for NEDD8 Activation and Conjugation
; FILE REFERENCE: 103576-127
; CURRENT APPLICATION NUMBER: US/09/216,430C
; CURRENT FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: PCT/US98/27141
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: US 60/068,209
; PRIOR FILING DATE: 1998-08-12
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 20

;
; LENGTH: 152
; TYPE: PRT
; ORGANISM: Human
; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (1)..(152)
; OTHER INFORMATION: UbcH8
US-09-216-430C-20

Query Match 60.3%; Score 41; DB 2; Length 152;
Best Local Similarity 63.6%; Pred. No. 9.5;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFK 11
:|||||:||||
Db 53 RISFPPEYFPK 63

RESULT 10
US-08-748-703-1
; Sequence 1, Application US/08748703
; Patent No. 5847094
; GENERAL INFORMATION:
; APPLICANT: Bandman, Olga
; APPLICANT: Goli, Surya K.
; TITLE OF INVENTION: NOVEL UBCH7-LIKE UBIQUITIN-CONJUGATING
; TITLE OF INVENTION: ENZYME
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
; STREET: 3174 Porter Drive
; CITY: Palo Alto
; STATE: CA
; COUNTRY: US
; ZIP: 94304
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/748,703
; FILING DATE: Herewith
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Billings, Lucy J.
; REGISTRATION NUMBER: 36,749
; REFERENCE/DOCKET NUMBER: PF-0161 US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-855-0555
; TELEFAX: 415-845-4166
; TELEX:
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 153 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; IMMEDIATE SOURCE:
; LIBRARY: Consensus
; CLONE: Consensus
US-08-748-703-1

Query Match 60.3%; Score 41; DB 1; Length 153;
Best Local Similarity 63.6%; Pred. No. 9.5;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFK 11
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Db 54 RISFPPEYFPK 64

RESULT 11
US-09-132-861-1
; Sequence 1, Application US/09132861
; Patent No. 6124123
; GENERAL INFORMATION:
; APPLICANT: Bandman, Olga
; APPLICANT: Goli, Surya K.
; TITLE OF INVENTION: NOVEL UBCH7-LIKE UBIQUITIN-CONJUGATING
; TITLE OF INVENTION: ENZYME
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
; STREET: 3174 Porter Drive
; CITY: Palo Alto
; STATE: CA
; COUNTRY: US
; ZIP: 94304
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/132,861
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/748,703
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Billings, Lucy J.
; REGISTRATION NUMBER: 36,749
; REFERENCE/DOCKET NUMBER: PF-0161 US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-855-0555
; TELEFAX: 415-845-4166
; TELEX:
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 153 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; IMMEDIATE SOURCE:
; LIBRARY: Consensus
; CLONE: Consensus
US-09-132-861-1

Query Match 60.3%; Score 41; DB 2; Length 153;
Best Local Similarity 63.6%; Pred. No. 9.5;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFK 11
:|||||:||||
Db 54 RISFPPEYFPK 64

RESULT 12
US-09-949-016-6465
; Sequence 6465, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768

; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6465
; LENGTH: 153
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-6465

Query Match 60.3%; Score 41; DB 2; Length 153;
Best Local Similarity 63.6%; Pred. No. 9.5;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
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DB 54 RISFPPEYFPK 64

RESULT 13
US-09-949-016-10584
; Sequence 10584, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10584
; LENGTH: 168
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-10584

Query Match 60.3%; Score 41; DB 2; Length 168;
Best Local Similarity 63.6%; Pred. No. 11;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
:|||||:||||
DB 69 RISFPPEYFPK 79

RESULT 14
US-09-331-568A-2
; Sequence 2, Application US/09331568A
; Patent No. 6570004
; GENERAL INFORMATION:
; APPLICANT: Martin J. Blaser
; APPLICANT: Mikio Karita
; TITLE OF INVENTION: dape GENE OF HELICOBACTER PYLORI AND
; FILE REFERENCE: dape MUTANT STRAINS OF HELICOBACTER PYLORI
; FILE REFERENCE: 22000.0072
; CURRENT APPLICATION NUMBER: US/09/331,568A
; CURRENT FILING DATE: 2001-06-18
; PRIOR APPLICATION NUMBER: PCT/US97/24147
; PRIOR FILING DATE: 1997-12-23
; PRIOR APPLICATION NUMBER: 60/033,824
; PRIOR FILING DATE: 1996-12-23
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 2

; LENGTH: 621
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:\No. 6570004e =
; OTHER INFORMATION: synthetic construct
US-09-331-568A-2

Query Match 60.3%; Score 41; DB 2; Length 621;
Best Local Similarity 63.6%; Pred. No. 46;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
:|||||:||||
DB 563 KVSIPKGFVFK 573

RESULT 15
US-09-331-568A-26
; Sequence 26, Application US/09331568A
; Patent No. 6570004
; GENERAL INFORMATION:
; APPLICANT: Martin J. Blaser
; APPLICANT: Mikio Karita
; TITLE OF INVENTION: dape GENE OF HELICOBACTER PYLORI AND
; FILE REFERENCE: dape MUTANT STRAINS OF HELICOBACTER PYLORI
; CURRENT APPLICATION NUMBER: US/09/331,568A
; CURRENT FILING DATE: 2001-06-18
; PRIOR APPLICATION NUMBER: PCT/US97/24147
; PRIOR FILING DATE: 1997-12-23
; PRIOR APPLICATION NUMBER: 60/033,824
; PRIOR FILING DATE: 1996-12-23
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 26
; LENGTH: 621
; TYPE: PRT
; ORGANISM: H. pylori
US-09-331-568A-26

Query Match 60.3%; Score 41; DB 2; Length 621;
Best Local Similarity 63.6%; Pred. No. 46;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
:|||||:||||
DB 563 KVSIPKGFVFK 573

Search completed: January 28, 2006, 06:24:31
Job time : 19.6667 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: January 28, 2006, 06:23:49 ; Search time 12.3333 Seconds
(without alignments)
10.536 Million cell updates/sec

Title: US-09-897-042-20
Perfect score: 64
Sequence: 1 TIKKISFPEGFP 12

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 75621 seqs, 10829074 residues

Total number of hits satisfying chosen parameters: 75621

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA New:*

- 1: /cgn2_6/prodata/2/pubpaa/US08_NEW_PUB.psp.*
- 2: /cgn2_6/prodata/2/pubpaa/US06_NEW_PUB.psp.*
- 3: /cgn2_6/prodata/2/pubpaa/US07_NEW_PUB.psp.*
- 4: /cgn2_6/prodata/2/pubpaa/PCT_NEW_PUB.psp.*
- 5: /cgn2_6/prodata/2/pubpaa/US09_NEW_PUB.psp.*
- 6: /cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.psp.*
- 7: /cgn2_6/prodata/2/pubpaa/US11_NEW_PUB.psp.*
- 8: /cgn2_6/prodata/2/pubpaa/US60_NEW_PUB.psp.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Score	Length	ID	Description
1	64	100.0	172	US-11-102-883-18	Sequence 18, Appl
2	64	100.0	289	US-11-102-883-26	Sequence 26, Appl
3	64	100.0	300	US-11-102-883-6	Sequence 6, Appl
4	46	71.9	225	US-10-209-208-1	Sequence 1, Appl
5	46	71.9	225	US-10-209-208-4	Sequence 4, Appl
6	46	71.9	225	US-10-209-208-20	Sequence 20, Appl
7	46	71.9	225	US-10-209-208-24	Sequence 24, Appl
8	46	71.9	225	US-11-218-880-1	Sequence 1, Appl
9	46	71.9	225	US-11-218-880-4	Sequence 4, Appl
10	46	71.9	225	US-11-218-880-20	Sequence 20, Appl
11	46	71.9	225	US-11-218-880-24	Sequence 24, Appl
12	46	71.9	225	US-11-100-988-2	Sequence 2, Appl
13	46	71.9	225	US-11-100-988-3	Sequence 3, Appl
14	46	71.9	226	US-10-209-208-6	Sequence 6, Appl
15	46	71.9	226	US-11-218-880-6	Sequence 6, Appl
16	41	64.1	225	US-10-209-208-8	Sequence 8, Appl
17	41	64.1	225	US-10-209-208-79	Sequence 79, Appl
18	41	64.1	225	US-11-218-880-8	Sequence 8, Appl
19	41	64.1	225	US-11-100-988-5	Sequence 5, Appl
20	40	62.5	343	US-10-131-826A-162	Sequence 162, App
21	36	56.2	3567	US-10-453-372-1112	Sequence 112, App
22	35	54.7	125	US-09-978-360A-442	Sequence 442, App
23	35	54.7	198	US-10-131-826A-550	Sequence 550, App
24	35	54.7	495	US-10-467-962B-93	Sequence 93, Appl
25	35	54.7	661	US-11-019-711-107	Sequence 107, App

26	35	54.7	964	7	US-11-016-706-39	Sequence 39, Appl
27	34	53.1	233	7	US-11-179-411-27	Sequence 27, Appl
28	34	53.1	233	7	US-11-179-411-33	Sequence 33, Appl
29	34	53.1	233	7	US-11-175-766-27	Sequence 27, Appl
30	34	53.1	233	7	US-11-175-766-33	Sequence 33, Appl
31	34	53.1	291	7	US-11-010-239-119	Sequence 119, App
32	33	51.6	283	7	US-11-102-883-32	Sequence 32, Appl
33	33	51.6	294	7	US-11-102-883-30	Sequence 30, Appl
34	33	51.6	833	6	US-10-467-657-3876	Sequence 3876, Ap
35	33	51.6	1015	6	US-10-957-569-51	Sequence 51, Appl
36	33	51.6	3483	6	US-10-453-372-40	Sequence 40, Appl
37	33	51.6	3546	6	US-10-453-372-32	Sequence 32, Appl
38	32.5	50.8	626	6	US-10-467-657-6426	Sequence 6426, Ap
39	32.5	50.8	626	6	US-10-467-657-7618	Sequence 7618, Ap
40	32	50.0	34	6	US-10-467-657-4010	Sequence 4010, Ap
41	32	50.0	119	7	US-11-073-605-12	Sequence 12, Appl
42	32	50.0	221	7	US-11-076-164-16	Sequence 16, Appl
43	32	50.0	251	6	US-10-485-517-311	Sequence 311, App
44	32	50.0	258	6	US-10-793-626-2360	Sequence 2360, Ap
45	32	50.0	278	7	US-11-165-067A-5	Sequence 5, Appl

ALIGNMENTS

RESULT 1

US-11-102-883-18
; Sequence 18, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lampung, Norbert
; APPLICANT: Crameri, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods and
; TITLE OF INVENTION: Uses Thereof
; FILE REFERENCE: 03100234pa
; CURRENT APPLICATION NUMBER: US/11/102,883
; CURRENT FILING DATE: 2005-04-11
; PRIOR APPLICATION NUMBER: EP02022774.0
; PRIOR FILING DATE: 2002-10-11
; PRIOR APPLICATION NUMBER: PCT/EP2003/011190
; PRIOR FILING DATE: 2003-10-09
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: Patentin version 3.2
; SEQ ID NO 18
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Betula verrucosa
US-11-102-883-18

Query Match 100.0%; Score 64; DB 7; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.00011;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12

Db 65 TIKKISFPEGFP 76

RESULT 2

US-11-102-883-26
; Sequence 26, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lampung, Norbert
; APPLICANT: Crameri, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods and
; TITLE OF INVENTION: Uses Thereof

;; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
;; FILE REFERENCE: UC083.LCP2CP2
;; CURRENT APPLICATION NUMBER: US/10/209,208
;; CURRENT FILING DATE: 2002-07-29
;; PRIOR APPLICATION NUMBER: 10/121,258
;; PRIOR FILING DATE: 2002-04-10
;; PRIOR APPLICATION NUMBER: 09/866,538
;; PRIOR FILING DATE: 2001-05-24
;; PRIOR APPLICATION NUMBER: 09/794,308
;; PRIOR FILING DATE: 2001-02-26
;; NUMBER OF SEQ ID NOS: 80
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 20
;; LENGTH: 225
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: DsRed with I125R
US-10-209-208-20

Query Match 71.9%; Score 46; DB 6; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 7

US-10-209-208-24
;; Sequence 24, Application US/10209208
;; Publication No. US20050244921A1
;; GENERAL INFORMATION:
;; APPLICANT: Tsien, Roger
;; APPLICANT: Campbell, Robert
;; APPLICANT: Geoffrey Baird

;; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
;; FILE REFERENCE: UC083.LCP2CP2
;; CURRENT APPLICATION NUMBER: US/10/209,208
;; CURRENT FILING DATE: 2002-07-29
;; PRIOR APPLICATION NUMBER: 10/121,258
;; PRIOR FILING DATE: 2002-04-10
;; PRIOR APPLICATION NUMBER: 09/866,538
;; PRIOR FILING DATE: 2001-05-24
;; PRIOR APPLICATION NUMBER: 09/794,308
;; PRIOR FILING DATE: 2001-02-26
;; NUMBER OF SEQ ID NOS: 80
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 24
;; LENGTH: 225
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: DsRed polypeptide variant "T1" with I125R mutation
US-10-209-208-24

Query Match 71.9%; Score 46; DB 6; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 8

US-11-218-880-1
;; Sequence 1, Application US/11218880
;; Publication No. US20060003420A1
;; GENERAL INFORMATION:
;; APPLICANT: Tsien, Roger
;; APPLICANT: Campbell, Robert
;; APPLICANT: Geoffrey Baird

;; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
;; FILE REFERENCE: UC083.LCP2CP1
;; CURRENT APPLICATION NUMBER: US/11/218,880
;; CURRENT FILING DATE: 2005-09-01
;; PRIOR APPLICATION NUMBER: US/10/121,258
;; PRIOR FILING DATE: 2002-04-10
;; PRIOR APPLICATION NUMBER: 09/794,308
;; PRIOR FILING DATE: 2001-02-26
;; PRIOR APPLICATION NUMBER: 09/866,538
;; PRIOR FILING DATE: 2001-05-24
;; NUMBER OF SEQ ID NOS: 78
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 1
;; LENGTH: 225
;; TYPE: PRT
;; ORGANISM: Discosoma sp.
;; FEATURE:
;; NAME/KEY: misc.feature
;; LOCATION: (1)...(225)
;; OTHER INFORMATION: wild-type DsRed
US-11-218-880-1

Query Match 71.9%; Score 46; DB 7; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 9

US-11-218-880-4
;; Sequence 4, Application US/11218880
;; Publication No. US20060003420A1
;; GENERAL INFORMATION:
;; APPLICANT: Tsien, Roger
;; APPLICANT: Campbell, Robert
;; APPLICANT: Geoffrey Baird

;; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
;; FILE REFERENCE: UC083.LCP2CP1
;; CURRENT APPLICATION NUMBER: US/11/218,880
;; CURRENT FILING DATE: 2005-09-01
;; PRIOR APPLICATION NUMBER: US/10/121,258
;; PRIOR FILING DATE: 2002-04-10
;; PRIOR APPLICATION NUMBER: 09/794,308
;; PRIOR FILING DATE: 2001-02-26
;; PRIOR APPLICATION NUMBER: 09/866,538
;; PRIOR FILING DATE: 2001-05-24
;; NUMBER OF SEQ ID NOS: 78
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 4
;; LENGTH: 225
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: DsRed polypeptide variant "T1"
US-11-218-880-4

Query Match 71.9%; Score 46; DB 7; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 10

US-11-218-880-20

; Sequence 20, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 20
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed with I125R
US-11-218-880-20

Query Match 71.9%; Score 46; DB 7; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 11

US-11-218-880-24
; Sequence 24, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 24
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed polypeptide variant "T1" with I125R mutation
US-11-218-880-24

Query Match 71.9%; Score 46; DB 7; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 12

US-11-100-988-2
; Sequence 2, Application US/11100988
; Publication No. US20060008878A1
; GENERAL INFORMATION:
; APPLICANT: Bevis, Brooke
; TITLE OF INVENTION: MONOMERIC RED FLUORESCENT PROTEINS
; FILE REFERENCE: 092234-9002-US01
; CURRENT APPLICATION NUMBER: US/11/100,988
; CURRENT FILING DATE: 2005-04-07
; PRIOR APPLICATION NUMBER: US 60/560,340
; PRIOR FILING DATE: 2004-04-07
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Unknown
; FEATURE:
; OTHER INFORMATION: wild-type red fluorescent protein based on Discosoma species
US-11-100-988-2

Query Match 71.9%; Score 46; DB 7; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 13

US-11-100-988-3
; Sequence 3, Application US/11100988
; Publication No. US20060008878A1
; GENERAL INFORMATION:
; APPLICANT: Bevis, Brooke
; TITLE OF INVENTION: MONOMERIC RED FLUORESCENT PROTEINS
; FILE REFERENCE: 092234-9002-US01
; CURRENT APPLICATION NUMBER: US/11/100,988
; CURRENT FILING DATE: 2005-04-07
; PRIOR APPLICATION NUMBER: US 60/560,340
; PRIOR FILING DATE: 2004-04-07
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 3
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Unknown
; FEATURE:
; OTHER INFORMATION: rapidly maturing DaRed.T4 red fluorescent protein based on
; OTHER INFORMATION: Discosoma species
US-11-100-988-3

Query Match 71.9%; Score 46; DB 7; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 14
US-10-209-208-6
; Sequence 6, Application US/10209208

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; Publication No. US2005024921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; TITLE OF INVENTION: FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6
; LENGTH: 226
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed polypeptide variant "dimer2"
US-10-209-208-6

Query Match          71.9%; Score 46; DB 6; Length 226;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY      3 KKISFPEGF 11
Db      84 KKLSFPEGF 92

RESULT 15
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; Sequence 6, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; TITLE OF INVENTION: PROTEIN VARIANTS AND METHODS FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6
; LENGTH: 226
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed polypeptide variant "dimer2"
US-11-218-880-6

Query Match          71.9%; Score 46; DB 7; Length 226;
Best Local Similarity 88.9%; Pred. No. 0.24;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
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QY      3 KKISFPEGF 11
Db      84 KKLSFPEGF 92
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OM protein - protein search, using sw model

Run on: January 28, 2006, 06:23:08 ; Search time 79.6667 Seconds
(without alignments)
62.937 Million cell updates/sec

Title: US-09-897-042-20
Perfect score: 64
Sequence: 1 TIKKISFPEGFP 12

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

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Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
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2	64	100.0	31	4	Sequence 20, Appli
3	64	100.0	125	5	Sequence 3, Appli
4	64	100.0	159	3	Sequence 5, Appli
5	64	100.0	159	3	Sequence 1, Appli
6	64	100.0	159	3	Sequence 34, Appli
7	64	100.0	159	3	Sequence 38, Appli
8	64	100.0	159	3	Sequence 40, Appli
9	64	100.0	159	3	Sequence 6, Appli
10	64	100.0	159	4	Sequence 47, Appli
11	64	100.0	160	4	Sequence 2, Appli
12	64	100.0	160	4	Sequence 3, Appli
13	64	100.0	160	4	Sequence 4, Appli
14	64	100.0	160	4	Sequence 5, Appli
15	64	100.0	160	4	Sequence 6, Appli
16	64	100.0	160	4	Sequence 7, Appli
17	64	100.0	160	5	Sequence 8, Appli
18	64	100.0	160	5	Sequence 7, Appli
19	64	100.0	161	4	Sequence 44, Appli
20	64	100.0	161	4	Sequence 45, Appli
21	64	100.0	195	5	Sequence 18, Appli
22	64	100.0	195	5	Sequence 20, Appli
23	61	95.3	159	3	Sequence 36, Appli
24	61	95.3	159	3	Sequence 39, Appli
25	61	95.3	159	3	Sequence 42, Appli
26	58	90.6	159	3	Sequence 37, Appli
27	58	90.6	159	4	Sequence 2, Appli

28	58	90.6	159	4	US-10-001-245-3	Sequence 3, Appli
29	58	90.6	159	4	US-10-001-245-4	Sequence 4, Appli
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31	58	90.6	159	4	US-10-001-245-9	Sequence 9, Appli
32	58	90.6	159	4	US-10-001-245-92	Sequence 92, Appli
33	58	90.6	159	4	US-10-719-553-37	Sequence 37, Appli
34	58	90.6	159	4	US-10-698-855-5	Sequence 5, Appli
35	58	90.6	160	4	US-10-001-245-1	Sequence 1, Appli
36	58	90.6	160	4	US-10-001-245-6	Sequence 6, Appli
37	58	90.6	160	4	US-10-001-245-7	Sequence 7, Appli
38	58	90.6	160	4	US-10-001-245-8	Sequence 8, Appli
39	58	90.6	160	4	US-10-001-245-10	Sequence 10, Appli
40	58	90.6	160	4	US-10-001-245-11	Sequence 11, Appli
41	58	90.6	160	4	US-10-001-245-12	Sequence 12, Appli
42	58	90.6	160	4	US-10-440-516-1	Sequence 1, Appli
43	58	90.6	160	4	US-10-440-516-11	Sequence 11, Appli
44	58	90.6	160	4	US-10-440-516-12	Sequence 12, Appli
45	58	90.6	160	4	US-10-440-516-15	Sequence 15, Appli

ALIGNMENTS

RESULT 1
US-09-897-042-20
; Sequence 20, Application US/09897042
; Patent No. US20020018779A1
; GENERAL INFORMATION:
; APPLICANT: VALENTA, Rudolf
; APPLICANT: VRTALA, Susanne
; APPLICANT: VANGELISTA, Luca
; APPLICANT: EICHLER, Hans-Georg
; APPLICANT: SPERR, Wolfgang R.
; APPLICANT: VALENT, Peter
; APPLICANT: EBNER, Christof
; APPLICANT: KRAFT, Dietrich
; APPLICANT: GRONLUND, Hans
; TITLE OF INVENTION: NON-ANAPHYLACTIC FORMS OF ALLERGENS AND THEIR USE
; FILE REFERENCE: 1614-251P
; CURRENT APPLICATION NUMBER: US/09/897,042
; CURRENT FILING DATE: 2001-07-07
; NUMBER OF SEQ ID NOS: 21
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 20
; LENGTH: 12
; TYPE: PRT
; ORGANISM: Betula sp.
US-09-897-042-20

Query Match 100.0%; Score 64; DB 3; Length 12;
Best Local Similarity 100.0%; Pred. No. 0.00031;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TIKKISFPEGFP 12
| | | | | | | | | | | |
Db 1 TIKKISFPEGFP 12

RESULT 2
US-10-026-911-3
; Sequence 3, Application US/10026911
; Publication No. US20030078201A1
; GENERAL INFORMATION:
; APPLICANT: Focke, Margarete
; APPLICANT: Mahler, Vera
; APPLICANT: Sperr, Wolfgang R.
; APPLICANT: Valent, Peter
; APPLICANT: Kraft, Dietrich
; APPLICANT: Valenta, Rudolf
; TITLE OF INVENTION: Allergy Vaccines and Their Preparation
; FILE REFERENCE: 0273-0005
; CURRENT APPLICATION NUMBER: US/10/026,911
; CURRENT FILING DATE: 2002-07-24

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; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 31
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:
; OTHER INFORMATION: solvent-exposed peptide
US-10-026-911-3

Query Match          100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.00086;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 5 TIKKISFPEGFP 16

RESULT 3
US-10-799-514-5
; Sequence 5, Application US/10799514
; Publication No. US20040241178A1
; GENERAL INFORMATION:
; APPLICANT: Spertini, Francois
; APPLICANT: Cortesey, Blaise
; TITLE OF INVENTION: Allergen Peptide Fragments and Use Thereof
; FILE REFERENCE: 25720-502
; CURRENT APPLICATION NUMBER: US/10/799,514
; PRIOR FILING DATE: 2004-03-12
; PRIOR APPLICATION NUMBER: 60/455,004
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 5
; LENGTH: 125
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: Peptide
US-10-799-514-5

Query Match          100.0%; Score 64; DB 5; Length 125;
Best Local Similarity 100.0%; Pred. No. 0.0038;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 53 TIKKISFPEGFP 64

RESULT 4
US-09-981-009B-1
; Sequence 1, Application US/09981009B
; Publication No. US20030041354A1
; GENERAL INFORMATION:
; APPLICANT: Kjaerulf, Soren
; APPLICANT: Roggen, Erwin
; TITLE OF INVENTION: Transgenic Plants
; FILE REFERENCE: 10082.200-US
; CURRENT APPLICATION NUMBER: US/09/981,009B
; CURRENT FILING DATE: 2002-06-14
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula pendula
US-09-981-009B-1

Query Match          100.0%; Score 64; DB 3; Length 159;
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Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 52 TIKKISFPEGFP 63

RESULT 5
US-09-847-208-34
; Sequence 34, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 34
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-34

Query Match          100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 52 TIKKISFPEGFP 63

RESULT 6
US-09-847-208-38
; Sequence 38, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 38
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-38

Query Match          100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 52 TIKKISFPEGFP 63

RESULT 7
US-09-847-208-40
; Sequence 40, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
```

; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: AGE-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 40
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-40

Query Match 100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 52 TIKKISFPEGFP 63

RESULT 8

US-09-957-806A-6
; Sequence 6, Application US/09957806A
; Publication No. US20050181446A1
; GENERAL INFORMATION:
; APPLICANT: Roggen, Erwin
; APPLICANT: Ernst, Steffen
; APPLICANT: Svendsen, Allan
; APPLICANT: Friis, Esben
; APPLICANT: Osten, Claus
; TITLE OF INVENTION: PROTEIN VARIANTS HAVING MODIFIED IMMUNOGENICITY
; FILE REFERENCE: 10021.204-US
; CURRENT APPLICATION NUMBER: US/09/957,806A
; CURRENT FILING DATE: 2001-09-21
; NUMBER OF SEQ ID NOS: 248
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 6
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula pendula
US-09-957-806A-6

Query Match 100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 52 TIKKISFPEGFP 63

RESULT 9

US-10-440-516-47
; Sequence 47, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 47
; LENGTH: 159

; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (1)-(159)
; OTHER INFORMATION: Point mutated Bet v 1 allergen (amino acid deletion of residue
; OTHER INFORMATION: G111)
US-10-440-516-47

Query Match 100.0%; Score 64; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 52 TIKKISFPEGFP 63

RESULT 10

US-10-440-516-2
; Sequence 2, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
US-10-440-516-2

Query Match 100.0%; Score 64; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
Db 53 TIKKISFPEGFP 64

RESULT 11

US-10-440-516-3
; Sequence 3, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 3
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (1)-(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-3

Query Match 100.0%; Score 64; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TIKKISFPEGPP 12
Db 52 TIKKISFPEGPP 63

RESULT 12
US-10-440-516-4
; Sequence 4, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 4
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; NAME/KEY: MISC FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-4

Query Match 100.0%; Score 64; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TIKKISFPEGPP 12
Db 52 TIKKISFPEGPP 63

RESULT 13
US-10-440-516-5
; Sequence 5, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 5
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; NAME/KEY: MISC FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-5

Query Match 100.0%; Score 64; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TIKKISFPEGPP 12

Db 52 TIKKISFPEGPP 63

RESULT 14
US-10-440-516-6
; Sequence 6, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 6
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; NAME/KEY: MISC FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-6

Query Match 100.0%; Score 64; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TIKKISFPEGPP 12
Db 52 TIKKISFPEGPP 63

RESULT 15
US-10-440-516-7
; Sequence 7, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 7
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; NAME/KEY: MISC FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-7

Query Match 100.0%; Score 64; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0049;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TIKKISFPEGPP 12
Db 52 TIKKISFPEGPP 63

Search completed: January 28, 2006, 06:43:58

Job time : 79.6667 secs



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OM protein - protein search, using sw model

Run on: January 28, 2006, 06:20:07 ; Search time 18.6667 Seconds
(without alignments)
53.149 Million cell updates/sec

Title: US-09-897-042-20
Perfect score: 64
Sequence: 1 TIKKISPEGFP 12

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents AA:*
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2: /cgn2_6/prodata/1/iaa/6 COMB.pep.*
3: /cgn2_6/prodata/1/iaa/H COMB.pep.*
4: /cgn2_6/prodata/1/iaa/ECTUS COMB.pep.*
5: /cgn2_6/prodata/1/iaa/RE COMB.pep.*
6: /cgn2_6/prodata/1/iaa/backfile1.pep.*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	64	100.0	160	1	US-07-847-010-23 Sequence 23, Appl
2	53	82.8	160	1	US-07-847-010-3 Sequence 3, Appli
3	46	71.9	225	2	US-09-866-538-12 Sequence 12, Appl
4	46	71.9	225	2	US-09-865-291-12 Sequence 12, Appl
5	46	71.9	240	2	US-10-152-296-2 Sequence 2, Appli
6	45	70.3	431	2	US-09-248-796A-14528 Sequence 14528, A
7	42	65.6	514	2	US-08-688-988-35 Sequence 35, Appl
8	41.5	64.8	98	2	US-09-205-658-194 Sequence 194, App
9	41.5	64.8	122	2	US-09-205-658-175 Sequence 175, App
10	41.5	64.8	632	2	US-09-205-658-160 Sequence 160, App
11	41.5	64.8	636	2	US-09-248-847-273 Sequence 273, App
12	40	62.5	414	2	US-08-823-516-145 Sequence 145, App
13	40	62.5	434	1	US-09-940-244-145 Sequence 145, App
14	40	62.5	434	2	US-09-381-212-145 Sequence 145, App
15	40	62.5	434	2	US-09-713-601A-145 Sequence 145, App
16	40	62.5	434	2	US-09-489-847-271 Sequence 271, App
17	40	62.5	443	2	US-09-489-847-132 Sequence 132, App
18	40	62.5	709	2	US-09-107-532A-5318 Sequence 5318, Ap
19	39	60.9	384	2	US-09-134-000C-5262 Sequence 5262, Ap
20	38	59.4	300	2	US-09-902-540-13351 Sequence 13351, A
21	37	57.8	247	2	US-09-248-796A-14142 Sequence 14142, A
22	37	57.8	318	2	US-09-705-534-4 Sequence 4, Appli
23	37	57.8	324	2	US-09-252-991A-26760 Sequence 26760, A
24	37	57.8	415	2	US-08-688-988-34 Sequence 34, Appl
25	37	57.8	524	2	US-08-688-988-10 Sequence 10, Appl
26	37	57.8	525	2	US-08-688-988-10 Sequence 10, Appl
27	37	57.8	793	1	US-08-468-558-5 Sequence 5, Appli

Sequence 5, Appli
Sequence 44, Appl
Sequence 2, Appl
Sequence 44, Appl
Sequence 37, Appl
Sequence 36, Appl
Sequence 7144, Ap
Sequence 1212, Ap
Sequence 611, App
Sequence 340, App
Sequence 8, Appli
Sequence 110, App
Sequence 1, Appli
Sequence 4, Appli
Sequence 2898, Ap
Sequence 5167, Ap
Sequence 5345, Ap

28 37 57.8 793 2 US-08-676-444-5
29 37 57.8 1751 2 US-09-136-574A-44
30 36 56.2 158 2 US-09-964-722-2
31 36 56.2 163 2 US-09-861-451A-44
32 36 56.2 310 1 US-08-129-456A-37
33 36 56.2 312 2 US-08-360-821B-36
34 36 56.2 312 2 US-09-851-026-36
35 36 56.2 320 2 US-09-949-016-7144
36 35 54.7 94 2 US-09-198-452A-1212
37 35 54.7 105 2 US-09-438-185A-611
38 35 54.7 111 2 US-09-149-476-340
39 35 54.7 125 2 US-09-599-360B-8
40 35 54.7 125 2 US-09-599-360B-110
41 35 54.7 154 1 US-08-363-010-1
42 35 54.7 154 1 US-08-911-434A-4
43 35 54.7 185 2 US-09-583-110-2898
44 35 54.7 196 2 US-09-107-433-5167
45 35 54.7 372 2 US-09-328-352-5345

ALIGNMENTS

RESULT 1
US-07-847-010-23
; Sequence 23, Application US/07847010
; Patent No. 5693495
; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikerstorfer, Arnold
; APPLICANT: Valentia, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
; APPLICANT: Rumpold, Helmut
; APPLICANT: Scheiner, Otto
; APPLICANT: Emer, Christof
; APPLICANT: Ferreira, Fatima
; TITLE OF INVENTION: Allergens of Alder Pollen and
; TITLE OF INVENTION: Applications Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 23:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:

; ORGANISM: birch (Betula sp.)
; IMMEDIATE SOURCE:
; LIBRARY: POLLEN FROM ALLERGO AB, ENGELHOLM, SWEDEN
US-07-847-010-23

Query Match 100.0%; Score 64; DB 1; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.00029;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
|||:|||||
Db 53 TIKKISFPEGFP 64

RESULT 2
US-07-847-010-3
; Sequence 3, Application US/07847010
; Patent No. 5693495

; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikerstorfer, Arnold
; APPLICANT: Valenta, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
; APPLICANT: Rumpold, Helmut
; APPLICANT: Scheiner, Otto
; APPLICANT: Ebner, Christof
; APPLICANT: Ferreira, Fatima
; TITLE OF INVENTION: Allergens of Alder Pollen and
; TITLE OF INVENTION: Applications Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711

; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C.
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; TELEX: 66141 PENNIE

; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHEetical: NO
; ORIGINAL SOURCE:
; ORGANISM: Alder (Alnus sp.)
US-07-847-010-3

Query Match 82.8%; Score 53; DB 1; Length 160;
Best Local Similarity 83.3%; Pred. No. 0.0033;
Matches 10; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 TIKKISFPEGFP 12
|||:|||||

Db 53 TIKKISFPEGSP 64

RESULT 3
US-09-866-538-12
; Sequence 12, Application US/09866538
; Patent No. 6852849
; GENERAL INFORMATION:
; APPLICANT: REGENTS OF THE UNIVERSITY OF CALIFORNIA
; APPLICANT: TSJEN, Roger
; APPLICANT: Campbell, Robert
; TITLE OF INVENTION: NON-OLIGOMERIZING FLUORESCENT PROTEINS
; FILE REFERENCE: REGEN1530-2
; CURRENT APPLICATION NUMBER: US/09/866,538
; CURRENT FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 29
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 12
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Discosoma sp.
US-09-866-538-12

Query Match 71.9%; Score 46; DB 2; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.97;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 4
US-09-865-291-12
; Sequence 12, Application US/09865291
; Patent No. 690304
; GENERAL INFORMATION:
; APPLICANT: REGENTS OF THE UNIVERSITY OF CALIFORNIA
; APPLICANT: TSJEN, Roger
; APPLICANT: TING, Alice
; APPLICANT: ZHANG, Jin
; TITLE OF INVENTION: EMISSION RATIO-METRIC INDICATORS OF PHOSPHORYLATION
; FILE REFERENCE: REGEN1550
; CURRENT APPLICATION NUMBER: US/09/865,291
; CURRENT FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 42
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 12
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Discosoma sp.
US-09-865-291-12

Query Match 71.9%; Score 46; DB 2; Length 225;
Best Local Similarity 88.9%; Pred. No. 0.97;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 KKISFPEGF 11
||:|||||
Db 83 KKLSFPEGF 91

RESULT 5
US-10-152-296-2
; Sequence 2, Application US/10152296
; Patent No. 6723537
; GENERAL INFORMATION:
; APPLICANT: Feelle, Beau
; APPLICANT: Rigel Pharmaceuticals, Incorporated
; TITLE OF INVENTION: Directed Evolution of Protein in Mammalian Cells
; FILE REFERENCE: 021044-000110US
; CURRENT APPLICATION NUMBER: US/10/152,296
; CURRENT FILING DATE: 2002-12-10

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; PRIOR APPLICATION NUMBER: US 60/291,871
; PRIOR FILING DATE: 2001-05-18
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 2
; LENGTH: 240
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:mammalian
; OTHER INFORMATION: codon-optimized variant (DERED) of Discosoma sp.
; OTHER INFORMATION: "red" red fluorescent protein (RFP)
US-10-152-296-2

Query Match          71.9%; Score 46; DB 2; Length 240;
Best Local Similarity 88.9%; Pred. No. 1;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY      3 KKISPEGF 11
Db      84 KKLSPPEGF 92

RESULT 6
US-09-248-796A-14528
; Sequence 14528, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; PRIOR FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 14528
; LENGTH: 431
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-14528

Query Match          70.3%; Score 45; DB 2; Length 431;
Best Local Similarity 66.7%; Pred. No. 3.1;
Matches 8; Conservative 2; Mismatches 2; Indels 2; Gaps 0;

QY      1 TIKKISFPEGFP 12
Db      117 TIKKTLPEGFP 128

RESULT 7
US-08-688-988-35
; Sequence 35, Application US/08688988B
; Patent No. 6096545
; GENERAL INFORMATION:
; APPLICANT: Lefebvre, Daniel D.
; APPLICANT: Malboobi, Mohammad A.
; TITLE OF INVENTION: PHOSPHATE STARVATION-INDUCIBLE PROTEINS
; FILE REFERENCE: PPL96-03
; CURRENT APPLICATION NUMBER: US/08/688,988B
; CURRENT FILING DATE: 1996-07-31
; NUMBER OF SEQ ID NOS: 48
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 35
; LENGTH: 514
; TYPE: PRT
; ORGANISM: Brassica napus
US-08-688-988-35
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Query Match          65.6%; Score 42; DB 2; Length 514;
Best Local Similarity 63.6%; Pred. No. 14;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY      1 TIKKISFPEGF 11
Db      32 TLRSASFPEGF 42

RESULT 8
US-09-205-658-194
; Sequence 194, Application US/09205658
; Patent No. 6861256
; GENERAL INFORMATION:
; APPLICANT: Ruvkun, Gary
; APPLICANT: Ogg, Scott
; TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
; TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
; FILE REFERENCE: 00786/351004
; CURRENT APPLICATION NUMBER: US/09/205,658
; CURRENT FILING DATE: 1998-12-03
; EARLIER APPLICATION NUMBER: 08/857,076
; EARLIER FILING DATE: 1997-05-15
; EARLIER APPLICATION NUMBER: 08/888,534
; EARLIER FILING DATE: 1997-07-07
; EARLIER APPLICATION NUMBER: US98/10080
; EARLIER FILING DATE: 1998-05-15
; NUMBER OF SEQ ID NOS: 328
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 194
; LENGTH: 98
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-205-658-194

Query Match          64.8%; Score 41.5; DB 2; Length 98;
Best Local Similarity 56.2%; Pred. No. 2.6;
Matches 9; Conservative 2; Mismatches 0; Indels 5; Gaps 1;

QY      2 IKKI-----SFPEGFP 12
Db      59 LKRIQLDFSFPEGFP 74

RESULT 9
US-09-205-658-175
; Sequence 175, Application US/09205658
; Patent No. 6861256
; GENERAL INFORMATION:
; APPLICANT: Ruvkun, Gary
; APPLICANT: Ogg, Scott
; TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
; TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
; FILE REFERENCE: 00786/351004
; CURRENT APPLICATION NUMBER: US/09/205,658
; CURRENT FILING DATE: 1998-12-03
; EARLIER APPLICATION NUMBER: 08/857,076
; EARLIER FILING DATE: 1997-05-15
; EARLIER APPLICATION NUMBER: 08/888,534
; EARLIER FILING DATE: 1997-07-07
; EARLIER APPLICATION NUMBER: US98/10080
; EARLIER FILING DATE: 1998-05-15
; NUMBER OF SEQ ID NOS: 328
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 175
; LENGTH: 122
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-205-658-175

Query Match          64.8%; Score 41.5; DB 2; Length 122;
Best Local Similarity 56.2%; Pred. No. 3.4;
Matches 9; Conservative 2; Mismatches 0; Indels 5; Gaps 1;
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QY 2 IKKI-----SPFEGFP 12
:|:| | | | | | | |
Db 29 LKRIQLDFSPFEGFP 44

RESULT 10
US-09-205-658-159
; Sequence 159, Application US/09205658
; Patent No. 6861256
; GENERAL INFORMATION:
; APPLICANT: Ruvkun, Gary
; APPLICANT: Ogg, Scott
; TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
; FILE REFERENCE: 00786/351004
; CURRENT APPLICATION NUMBER: US/09/205,658
; CURRENT FILING DATE: 1998-12-03
; EARLIER APPLICATION NUMBER: 08/857,076
; EARLIER FILING DATE: 1997-05-15
; EARLIER APPLICATION NUMBER: 08/888,534
; EARLIER FILING DATE: 1997-07-07
; EARLIER APPLICATION NUMBER: US98/10080
; EARLIER FILING DATE: 1998-05-15
; NUMBER OF SEQ ID NOS: 328
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 159
; LENGTH: 632
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-205-658-159

Query Match 64.8%; Score 41.5; DB 2; Length 632;
Best Local Similarity 56.2%; Pred. No. 21;
Matches 9; Conservative 2; Mismatches 0; Indels 5; Gaps 1;

QY 2 IKKI-----SPFEGFP 12
:|:| | | | | | | |
Db 317 LKRIQLDFSPFEGFP 332

RESULT 11
US-09-205-658-160
; Sequence 160, Application US/09205658
; Patent No. 6861256
; GENERAL INFORMATION:
; APPLICANT: Ruvkun, Gary
; APPLICANT: Ogg, Scott
; TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
; FILE REFERENCE: 00786/351004
; CURRENT APPLICATION NUMBER: US/09/205,658
; CURRENT FILING DATE: 1998-12-03
; EARLIER APPLICATION NUMBER: 08/857,076
; EARLIER FILING DATE: 1997-05-15
; EARLIER APPLICATION NUMBER: 08/888,534
; EARLIER FILING DATE: 1997-07-07
; EARLIER APPLICATION NUMBER: US98/10080
; EARLIER FILING DATE: 1998-05-15
; NUMBER OF SEQ ID NOS: 328
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 160
; LENGTH: 636
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-205-658-160

Query Match 64.8%; Score 41.5; DB 2; Length 636;
Best Local Similarity 56.2%; Pred. No. 21;
Matches 9; Conservative 2; Mismatches 0; Indels 5; Gaps 1;

QY 2 IKKI-----SPFEGFP 12
:|:| | | | | | | |

Db 319 LKRIQLDFSPFEGFP 334

RESULT 12
US-09-489-847-273
; Sequence 273, Application US/09489847
; Patent No. 6476195
; GENERAL INFORMATION:
; APPLICANT: Rosen et al
; TITLE OF INVENTION: 98 Human Secreted Proteins
; FILE REFERENCE: P2031P1
; CURRENT APPLICATION NUMBER: US/09/489,847
; CURRENT FILING DATE: 2000-01-24
; EARLIER APPLICATION NUMBER: PCT/US99/17130
; EARLIER FILING DATE: 1999-07-29
; EARLIER APPLICATION NUMBER: 60/094,657
; EARLIER FILING DATE: 1998-07-30
; EARLIER APPLICATION NUMBER: 60/095,486
; EARLIER FILING DATE: 1998-08-05
; EARLIER APPLICATION NUMBER: 60/096,319
; EARLIER FILING DATE: 1998-08-12
; EARLIER APPLICATION NUMBER: 60/095,454
; EARLIER FILING DATE: 1998-08-06
; EARLIER APPLICATION NUMBER: 60/095,455
; EARLIER FILING DATE: 1998-08-06
; NUMBER OF SEQ ID NOS: 376
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 273
; LENGTH: 414
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-489-847-273

Query Match 62.5%; Score 40; DB 2; Length 414;
Best Local Similarity 50.0%; Pred. No. 25;
Matches 6; Conservative 3; Mismatches 3; Indels 0; Gaps 0;

QY 1 TIKKISPFEGFP 12
|:|:| | | | | | | |
Db 234 TLMGANYFEGFP 245

RESULT 13
US-08-823-516-145
; Sequence 145, Application US/08823516
; Patent No. 5994069
; GENERAL INFORMATION:
; APPLICANT: Hall, Jeff G.
; APPLICANT: Lyamichev, Victor I.
; APPLICANT: Mast, Andrea L.
; APPLICANT: Brow, Mary Ann D.
; TITLE OF INVENTION: Detection Of Nucleic Acids By Multiple
; TITLE OF INVENTION: Sequential Invasive Cleavages
; NUMBER OF SEQUENCES: 163
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Medlen & Carroll, LLP
; STREET: 220 Montgomery Street, Suite 2200
; CITY: San Francisco
; STATE: California
; COUNTRY: United States Of America
; ZIP: 94104
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/823,516
; FILING DATE: 24-MAR-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/US97/01072
; FILING DATE: 21-JAN-1997

;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/759,038
;; FILING DATE: 02-DEC-1996
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/758,314
;; FILING DATE: 02-DEC-1996
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/756,386
;; FILING DATE: 29-NOV-1996
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/682,853
;; FILING DATE: 12-JUL-1996
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/599,491
;; FILING DATE: 24-JAN-1996
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Ingolia, Diane E.
;; REGISTRATION NUMBER: 40,027
;; REFERENCE/DOCKET NUMBER: FORS-02736
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (415) 705-8410
;; TELEFAX: (415) 397-8338
;; INFORMATION FOR SEQ ID NO: 145:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 434 amino acids
;; TYPE: amino acid
;; STRANDEDNESS: not relevant
;; TOPOLOGY: not relevant
;; MOLECULE TYPE: protein
;; US-08-823-516-145

Query Match 62.5%; Score 40; DB 1; Length 434;
Best Local Similarity 70.0%; Pred. No. 27;
Matches 7; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 3 KKISFPEGFP 12
||:|||||
Db 274 KKNFPPVGFP 283

RESULT 14
US-09-940-244-145
;; Sequence 145, Application US/09940244
;; Patent No. 6692917
;; GENERAL INFORMATION:
;; APPLICANT: Neri, Bruce P.
;; APPLICANT: Hall, Jeff G.
;; APPLICANT: Lyamichev, Victor
;; APPLICANT: Smith, Lloyd M.
;; TITLE OF INVENTION: Reactions on Dendrimers
;; FILE REFERENCE: FORS-06478
;; CURRENT APPLICATION NUMBER: US/09/940,244
;; CURRENT FILING DATE: 2002-05-06
;; NUMBER OF SEQ ID NOS: 422
;; SOFTWARE: Patentin version 3.1
;; SEQ ID NO 145
;; LENGTH: 434
;; TYPE: PRT
;; ORGANISM: Cunninghamella elegans
;; US-09-940-244-145

Query Match 62.5%; Score 40; DB 2; Length 434;
Best Local Similarity 70.0%; Pred. No. 27;
Matches 7; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 3 KKISFPEGFP 12
||:|||||
Db 274 KKNFPPVGFP 283

RESULT 15
US-09-381-212-145
;; Sequence 145, Application US/09381212

;; Patent No. 6872816
;; GENERAL INFORMATION:
;; APPLICANT: Hall, Jeff G.
;; Lyamichev, Victor I.
;; Mast, Andrea L.
;; Brown, Mary Ann D.
;; TITLE OF INVENTION: Detection Of Nucleic Acids By Multiple
;; Sequential Invasive Cleavages
;; NUMBER OF SEQUENCES: 190
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Medlen & Carroll, LLP
;; STREET: 220 Montgomery Street, Suite 2200
;; CITY: San Francisco
;; STATE: California
;; COUNTRY: United States Of America
;; ZIP: 94104
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Floppy disk
;; COMPUTER: IBM PC compatible
;; OPERATING SYSTEM: PC-DOS/MS-DOS
;; SOFTWARE: Patentin Release #1.0, Version #1.30
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/09/381,212
;; FILING DATE: 08-Feb-2000
;; CLASSIFICATION: <Unknown>
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: PCT/US97/01072
;; FILING DATE: 21-JAN-1997
;; APPLICATION NUMBER: US 08/759,038
;; FILING DATE: 02-DEC-1996
;; APPLICATION NUMBER: US 08/758,314
;; FILING DATE: 02-DEC-1996
;; APPLICATION NUMBER: US 08/756,386
;; FILING DATE: 29-NOV-1996
;; APPLICATION NUMBER: US 08/682,853
;; FILING DATE: 12-JUL-1996
;; APPLICATION NUMBER: US 08/599,491
;; FILING DATE: 24-JAN-1996
;; APPLICATION NUMBER: US 08/823,516
;; FILING DATE: 24-MAR-1997
;; ATTORNEY/AGENT INFORMATION:
;; NAME: MacKnight, Kamrin T.
;; REGISTRATION NUMBER: 38,230
;; REFERENCE/DOCKET NUMBER: FORS-03295
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (415) 705-8410
;; TELEFAX: (415) 397-8338
;; INFORMATION FOR SEQ ID NO: 145:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 434 amino acids
;; TYPE: amino acid
;; STRANDEDNESS: not relevant
;; TOPOLOGY: not relevant
;; MOLECULE TYPE: protein
;; SEQUENCE DESCRIPTION: SEQ ID NO: 145:
;; US-09-381-212-145

Query Match 62.5%; Score 40; DB 2; Length 434;
Best Local Similarity 70.0%; Pred. No. 27;
Matches 7; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 3 KKISFPEGFP 12
||:|||||
Db 274 KKNFPPVGFP 283

Search completed: January 28, 2006, 06:24:30
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OM protein - protein search, using sw model

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(without alignments)
10.536 Million cell updates/sec

Title: US-09-897-042-19
Perfect score: 64
Sequence: 1 GPGTIKKISFPE 12

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 75621 seqs, 10829074 residues

Total number of hits satisfying chosen parameters: 75621

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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7: /cgn2_6/ptodata/2/pubpaa/US11_NEW_PUB.pep.*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	64	100.0	172	7 US-11-102-883-18	Sequence 18, Appl
2	64	100.0	289	7 US-11-102-883-26	Sequence 26, Appl
3	64	100.0	300	7 US-11-102-883-6	Sequence 6, Appl
4	42	65.6	283	7 US-11-102-883-32	Sequence 32, Appl
5	42	65.6	294	7 US-11-102-883-30	Sequence 30, Appl
6	36	56.2	378	7 US-11-129-143-49	Sequence 49, Appl
7	36	56.2	608	7 US-11-226-701-8	Sequence 8, Appl
8	35	54.7	179	6 US-10-714-887-88	Sequence 88, Appl
9	35	54.7	285	6 US-10-821-234-983	Sequence 983, App
10	34	53.1	225	6 US-10-209-208-1	Sequence 1, Appl
11	34	53.1	225	6 US-10-209-208-4	Sequence 4, Appl
12	34	53.1	225	6 US-10-209-208-20	Sequence 20, Appl
13	34	53.1	225	6 US-10-209-208-24	Sequence 24, Appl
14	34	53.1	225	7 US-11-218-880-1	Sequence 1, Appl
15	34	53.1	225	7 US-11-218-880-4	Sequence 4, Appl
16	34	53.1	225	7 US-11-218-880-20	Sequence 20, Appl
17	34	53.1	225	7 US-11-218-880-24	Sequence 24, Appl
18	34	53.1	225	7 US-11-100-988-2	Sequence 2, Appl
19	34	53.1	225	7 US-11-100-988-3	Sequence 3, Appl
20	34	53.1	226	6 US-10-209-208-6	Sequence 6, Appl
21	34	53.1	226	7 US-11-218-880-6	Sequence 6, Appl
22	34	53.1	548	6 US-10-055-877-320	Sequence 320, App
23	34	53.1	674	6 US-10-055-877-319	Sequence 319, App
24	34	53.1	675	6 US-10-055-877-117	Sequence 117, App
25	34	53.1	675	6 US-10-055-877-317	Sequence 317, App

26	34	53.1	675	6 US-10-055-877-318	Sequence 318, App
27	34	53.1	768	7 US-11-147-109-10	Sequence 10, Appl
28	34	53.1	1436	7 US-11-052-554A-140	Sequence 140, App
29	33	51.6	186	7 US-10-793-626-3004	Sequence 3004, App
30	33	51.6	327	7 US-11-024-959-455	Sequence 455, App
31	33	51.6	664	6 US-10-624-932-10	Sequence 10, Appl
32	33	51.6	738	7 US-11-147-047-48	Sequence 48, Appl
33	33	51.6	795	7 US-11-052-554A-87	Sequence 87, Appl
34	33	51.6	869	6 US-10-453-372-50	Sequence 50, Appl
35	33	51.6	1024	7 US-11-145-631-2	Sequence 2, Appl
36	33	51.6	1070	7 US-11-147-047-49	Sequence 49, Appl
37	33	51.6	1204	7 US-11-145-631-5	Sequence 5, Appl
38	33	51.6	2612	6 US-10-453-372-38	Sequence 38, Appl
39	33	51.6	2669	6 US-10-453-372-36	Sequence 36, Appl
40	33	51.6	3104	6 US-10-453-372-34	Sequence 34, Appl
41	33	51.6	3104	6 US-10-453-372-62	Sequence 62, Appl
42	33	51.6	3104	6 US-10-453-372-64	Sequence 64, Appl
43	33	51.6	3130	6 US-10-453-372-42	Sequence 42, Appl
44	33	51.6	3483	6 US-10-453-372-40	Sequence 40, Appl
45	33	51.6	3546	6 US-10-453-372-32	Sequence 32, Appl

ALIGNMENTS

RESULT 1

US-11-102-883-18
; Sequence 18, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lampung, Norbert
; APPLICANT: Cramer, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods an
; TITLE OF INVENTION: Uses Thereof
; FILE REFERENCE: 03100234pa
; CURRENT APPLICATION NUMBER: US/11/102,883
; PRIOR FILING DATE: 2005-04-11
; PRIOR APPLICATION NUMBER: EP02022774.0
; PRIOR FILING DATE: 2002-10-11
; PRIOR APPLICATION NUMBER: PCT/EP2003/011190
; PRIOR FILING DATE: 2003-10-09
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: Patentin version 3.2
; SEQ ID NO 18
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Betula verrucosa
US-11-102-883-18

Query Match 100.0%; Score 64; DB 7; Length 172;
Best Local Similarity 100.0%; Pred. No. 5.5e-05;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12

Db 62 GPGTIKKISFPE 73

RESULT 2

US-11-102-883-26
; Sequence 26, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lampung, Norbert
; APPLICANT: Cramer, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods an
; TITLE OF INVENTION: Uses Thereof

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; FILE REFERENCE: 03100234pa
; CURRENT APPLICATION NUMBER: US/11/102,883
; CURRENT FILING DATE: 2005-04-11
; PRIOR APPLICATION NUMBER: EP02022774.0
; PRIOR FILING DATE: 2002-10-11
; PRIOR APPLICATION NUMBER: PCT/EP2003/011190
; PRIOR FILING DATE: 2003-10-09
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 26
; LENGTH: 289
; TYPE: PRT
; ORGANISM: tat-ii-bet v 1
US-11-102-883-26

Query Match      100.0%; Score 64; DB 7; Length 289;
Best Local Similarity 100.0%; Pred. No. 9.7e-05;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 GPGTIKKISFPE 12
      |||||
Db      179 GPGTIKKISFPE 190

RESULT 3
US-11-102-883-6
; Sequence 6, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lamping, Norbert
; APPLICANT: Cramer, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods and
; FILE REFERENCE: 03100234pa
; CURRENT APPLICATION NUMBER: US/11/102,883
; CURRENT FILING DATE: 2005-04-11
; PRIOR APPLICATION NUMBER: EP02022774.0
; PRIOR FILING DATE: 2002-10-11
; PRIOR APPLICATION NUMBER: PCT/EP2003/011190
; PRIOR FILING DATE: 2003-10-09
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 6
; LENGTH: 300
; TYPE: PRT
; ORGANISM: Human immunodeficiency virus + Homo sapiens + Betula verrucosa
US-11-102-883-6

Query Match      100.0%; Score 64; DB 7; Length 300;
Best Local Similarity 100.0%; Pred. No. 0.0001;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 GPGTIKKISFPE 12
      |||||
Db      190 GPGTIKKISFPE 201

RESULT 4
US-11-102-883-32
; Sequence 32, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lamping, Norbert
; APPLICANT: Cramer, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods and
; FILE REFERENCE: 03100234pa
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; CURRENT APPLICATION NUMBER: US/11/102,883
; CURRENT FILING DATE: 2005-04-11
; PRIOR APPLICATION NUMBER: EP02022774.0
; PRIOR FILING DATE: 2002-10-11
; PRIOR APPLICATION NUMBER: PCT/EP2003/011190
; PRIOR FILING DATE: 2003-10-09
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 32
; LENGTH: 283
; TYPE: PRT
; ORGANISM: tat-ii-api g 1
US-11-102-883-32

Query Match      65.6%; Score 42; DB 7; Length 283;
Best Local Similarity 58.3%; Pred. No. 1.1;
Matches 7; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY      1 GPGTIKKISFPE 12
      |||||
Db      178 GPGTIKKITLPD 189

RESULT 5
US-11-102-883-30
; Sequence 30, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lamping, Norbert
; APPLICANT: Cramer, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods and
; FILE REFERENCE: 03100234pa
; CURRENT APPLICATION NUMBER: US/11/102,883
; CURRENT FILING DATE: 2005-04-11
; PRIOR APPLICATION NUMBER: EP02022774.0
; PRIOR FILING DATE: 2002-10-11
; PRIOR APPLICATION NUMBER: PCT/EP2003/011190
; PRIOR FILING DATE: 2003-10-09
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 30
; LENGTH: 294
; TYPE: PRT
; ORGANISM: his-tat-ii-api g 1
US-11-102-883-30

Query Match      65.6%; Score 42; DB 7; Length 294;
Best Local Similarity 58.3%; Pred. No. 1.1;
Matches 7; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY      1 GPGTIKKISFPE 12
      |||||
Db      189 GPGTIKKITLPD 200

RESULT 6
US-11-129-143-49
; Sequence 49, Application US/11129143
; Publication No. US20050266518A1
; GENERAL INFORMATION:
; APPLICANT: BERRY, Alan
; APPLICANT: BRETZEL, Werner
; APPLICANT: HUMBELIN, Markus
; APPLICANT: LOPEZ-ULIBARRI, Rual
; APPLICANT: MAYER, Anne F.
; APPLICANT: YELISEV, Alexei A.
; TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
; FILE REFERENCE: C38435/121966
; CURRENT APPLICATION NUMBER: US/11/129,143
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; CURRENT FILING DATE: 2005-05-13
; NUMBER OF SEQ ID NOS: 197
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 49
; LENGTH: 378
; TYPE: PRT
; ORGANISM: Paracoccus sp. R114
US-11-129-143-49

Query Match 56.2%; Score 36; DB 7; Length 378;
Best Local Similarity 50.0%; Pred. No. 19;
Matches 6; Conservative 1; Mismatches 5; Indels 0; Gaps 0;

QY 1 GGTIKKISFPE 12
|||:|
Db 209 GPGSISSFDPLE 220

RESULT 7
US-11-226-701-8
; Sequence 8, Application US/11226701
; Publication No. US20060009632A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals, Inc.
; APPLICANT: Kapeller-Libermann, Rosana
; APPLICANT: Robison, Keith E.
; APPLICANT: White, David
; APPLICANT: Williamson, Mark W.
; APPLICANT: Cook, William James
; APPLICANT: Meyers, Rachel E.
; APPLICANT: MacBeth, Kyle J.
; APPLICANT: Carroll, Joseph M.
; APPLICANT: Chun, Miyoung
; TITLE OF INVENTION: NOVEL 27875, 22025, 27420, 17906, 16319.
; FILE REFERENCE: 55092 AND 10218 MOLECULES AND USES THEREFOR
; CURRENT APPLICATION NUMBER: US/11/226,701
; CURRENT FILING DATE: 2005-09-14
; PRIOR APPLICATION NUMBER: US/10/386,414
; PRIOR FILING DATE: 2003-03-11
; PRIOR APPLICATION NUMBER: 09/426,282
; PRIOR FILING DATE: 1999-10-25
; PRIOR APPLICATION NUMBER: 09/668,266
; PRIOR FILING DATE: 2000-09-22
; PRIOR APPLICATION NUMBER: 09/330,970
; PRIOR FILING DATE: 1999-06-11
; PRIOR APPLICATION NUMBER: 09/724,599
; PRIOR FILING DATE: 2000-11-28
; PRIOR APPLICATION NUMBER: 09/860,193
; PRIOR FILING DATE: 2001-05-16
; PRIOR APPLICATION NUMBER: 09/571,689
; PRIOR FILING DATE: 2000-05-16
; PRIOR APPLICATION NUMBER: 10/283,023
; PRIOR FILING DATE: 2002-10-29
; PRIOR APPLICATION NUMBER: 60/335,044
; PRIOR FILING DATE: 2001-10-31
; PRIOR APPLICATION NUMBER: 10/010,943
; PRIOR FILING DATE: 2001-12-06
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 608
; TYPE: PRT
; ORGANISM: Homo Sapiens
; FEATURE:
; NAME/KEY: VARIANT
; LOCATION: (1)...(608)
; OTHER INFORMATION: Xaa = Any Amino Acid
US-11-226-701-8

Query Match 56.2%; Score 36; DB 7; Length 608;
Best Local Similarity 45.5%; Pred. No. 32;

Matches 5; Conservative 4; Mismatches 2; Indels 0; Gaps 0;
QY 2 PGTIKKISFPE 12
|||:|
Db 239 PCKVEEVSLE 249

RESULT 8
US-10-714-887-88
; Sequence 88, Application US/10714887
; Publication No. US20060015972A1
; GENERAL INFORMATION:
; APPLICANT: Mendel Biotechnology, Inc.
; APPLICANT: HEARD, Jacqueline
; APPLICANT: RIECHMANN, Jose Luis
; APPLICANT: CREELMAN, Robert
; APPLICANT: RATCLIFFE, Oliver
; APPLICANT: CANALES, Roger
; APPLICANT: REPETTI, Peter
; APPLICANT: KUMIMOTO, Roderick W
; APPLICANT: GUTTERSON, Neal
; APPLICANT: REUBER, T. Lynne
; APPLICANT: PINEDA, Omaira
; APPLICANT: SHERMAN, Bradley K
; TITLE OF INVENTION: PLANT TRANSCRIPTIONAL REGULATORS OF DROUGHT STRESS
; FILE REFERENCE: MBI0058-CIP
; CURRENT APPLICATION NUMBER: US/10/714,887
; CURRENT FILING DATE: 2003-11-13
; PRIOR APPLICATION NUMBER: 10/412,699
; PRIOR FILING DATE: 2003-04-10
; PRIOR APPLICATION NUMBER: 09/506,720
; PRIOR FILING DATE: 2000-02-17
; PRIOR APPLICATION NUMBER: 60/135,134
; PRIOR FILING DATE: 1999-05-20
; PRIOR APPLICATION NUMBER: 09/394,519
; PRIOR FILING DATE: 1999-09-13
; PRIOR APPLICATION NUMBER: 09/533,392
; PRIOR FILING DATE: 2000-03-22
; PRIOR APPLICATION NUMBER: 09/533,029
; PRIOR FILING DATE: 2000-03-22
; PRIOR APPLICATION NUMBER: 09/532,591
; PRIOR FILING DATE: 2000-03-22
; PRIOR APPLICATION NUMBER: 09/533,030
; PRIOR FILING DATE: 2000-03-22
; PRIOR APPLICATION NUMBER: 60/125,814
; PRIOR FILING DATE: 1999-03-23
; PRIOR APPLICATION NUMBER: 09/713,994
; PRIOR FILING DATE: 2000-11-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 430
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 88
; LENGTH: 179
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: G3643 polypeptide Orthologous to G47
US-10-714-887-88

Query Match 54.7%; Score 35; DB 6; Length 179;
Best Local Similarity 45.5%; Pred. No. 13;
Matches 5; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

QY 2 PGTIKKISFPE 12
|||:|
Db 66 PSSLDKLNFPF 76

RESULT 9
US-10-821-234-983
; Sequence 983, Application US/10821234
; Publication No. US2005025114A1
; GENERAL INFORMATION:

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; APPLICANT: Labat, Ivan
; APPLICANT: Stache-Crain, Birgit
; APPLICANT: Andarmani, Susan
; APPLICANT: Tang, Y. Tom
; TITLE OF INVENTION: Methods for Diagnosis and Treatment of Presclampsia
; FILE REFERENCE: 821A
; CURRENT APPLICATION NUMBER: US/10/821,234
; CURRENT FILING DATE: 2004-04-07
; PRIOR APPLICATION NUMBER: US 60/462,047
; PRIOR FILING DATE: 2003-04-07
; NUMBER OF SEQ ID NOS: 1704
; SOFTWARE: pt_seq_genes Version 1.0
; SEQ ID NO 983
; LENGTH: 285
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-821-234-983

Query Match          54.7%; Score 35; DB 6; Length 285;
Best Local Similarity 54.5%; Pred. No. 22;
Matches 6; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

QY 1 GPGTIKKISFP 11
Db 31 GPSTLRPPSFP 41

RESULT 10
US-10-209-208-1
; Sequence 1, Application US/10209208
; Publication No. US20050244921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Discosoma sp.
; NAME/KEY: misc.feature
; LOCATION: (1)...(225)
; OTHER INFORMATION: wild-type DsRed
US-10-209-208-1

Query Match          53.1%; Score 34; DB 6; Length 225;
Best Local Similarity 85.7%; Pred. No. 26;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 6 KKISFPE 12
Db 83 KKLSFPE 89

RESULT 11
US-10-209-208-4
; Sequence 4, Application US/10209208
; Publication No. US20050244921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger

```

```

; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed polypeptide variant "T1"
US-10-209-208-4

Query Match          53.1%; Score 34; DB 6; Length 225;
Best Local Similarity 85.7%; Pred. No. 26;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 6 KKISFPE 12
Db 83 KKLSFPE 89

RESULT 12
US-10-209-208-20
; Sequence 20, Application US/10209208
; Publication No. US20050244921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 20
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed with I125R
US-10-209-208-20

Query Match          53.1%; Score 34; DB 6; Length 225;
Best Local Similarity 85.7%; Pred. No. 26;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 6 KKISFPE 12
Db 83 KKLSFPE 89

RESULT 13
US-10-209-208-24
; Sequence 24, Application US/10209208

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; Publication No. US20050244921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; TITLE OF INVENTION: FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 24
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed polypeptide variant "T1" with I125R mutation
US-10-209-208-24

Query Match          53.1%; Score 34; DB 6; Length 225;
Best Local Similarity 85.7%; Pred. No. 26;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy      6 KKISFPE 12
Db      83 KKLSFPE 89
      ||:||||

RESULT 14
US-11-218-880-1
; Sequence 1, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; TITLE OF INVENTION: PROTEIN VARIANTS AND METHODS FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Discosoma sp.
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(225)
; OTHER INFORMATION: wild-type DsRed
US-11-218-880-1

Query Match          53.1%; Score 34; DB 7; Length 225;
Best Local Similarity 85.7%; Pred. No. 26;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy      6 KKISFPE 12
Db      83 KKLSFPE 89
      ||:||||

; Publication No. US20050244921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; TITLE OF INVENTION: FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 24
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed polypeptide variant "T1" with I125R mutation
US-10-209-208-24

Query Match          53.1%; Score 34; DB 6; Length 225;
Best Local Similarity 85.7%; Pred. No. 26;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy      6 KKISFPE 12
Db      83 KKLSFPE 89
      ||:||||

RESULT 15
US-11-218-880-4
; Sequence 4, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; TITLE OF INVENTION: PROTEIN VARIANTS AND METHODS FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed polypeptide variant "T1"
US-11-218-880-4

Query Match          53.1%; Score 34; DB 7; Length 225;
Best Local Similarity 85.7%; Pred. No. 26;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy      6 KKISFPE 12
Db      83 KKLSFPE 89
      ||:||||

Search completed: January 28, 2006, 06:44:47
Job time : 13.3333 secs
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OM protein - protein search, using sw model

Run on: January 28, 2006, 06:23:08 ; Search time 79.6667 Seconds
(without alignments)
62.937 Million cell updates/sec

Title: US-09-897-042-19

Perfect score: 64

Sequence: 1 GPGTIKKISFPE 12

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Published Applications AA Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	ID	Description
1	64	100.0	12	3	US-09-897-042-19
2	64	100.0	31	4	US-10-026-911-3
3	64	100.0	125	5	US-10-799-514-5
4	64	100.0	159	3	US-09-981-009B-1
5	64	100.0	159	3	US-09-847-208-34
6	64	100.0	159	3	US-09-847-208-37
7	64	100.0	159	3	US-09-847-208-38
8	64	100.0	159	3	US-09-847-208-40
9	64	100.0	159	3	US-09-957-806A-6
10	64	100.0	159	4	US-10-001-245-2
11	64	100.0	159	4	US-10-001-245-3
12	64	100.0	159	4	US-10-001-245-4
13	64	100.0	159	4	US-10-001-245-5
14	64	100.0	159	4	US-10-001-245-9
15	64	100.0	159	4	US-10-001-245-92
16	64	100.0	159	4	US-10-440-516-47
17	64	100.0	159	4	US-10-719-553-37
18	64	100.0	159	4	US-10-698-855-5
19	64	100.0	160	4	US-10-001-245-1
20	64	100.0	160	4	US-10-001-245-6
21	64	100.0	160	4	US-10-001-245-7
22	64	100.0	160	4	US-10-001-245-8
23	64	100.0	160	4	US-10-001-245-10
24	64	100.0	160	4	US-10-001-245-11
25	64	100.0	160	4	US-10-001-245-12
26	64	100.0	160	4	US-10-440-516-1
27	64	100.0	160	4	US-10-440-516-2

28	64	100.0	160	4	US-10-440-516-3	Sequence 3, Appli
29	64	100.0	160	4	US-10-440-516-4	Sequence 4, Appli
30	64	100.0	160	4	US-10-440-516-5	Sequence 5, Appli
31	64	100.0	160	4	US-10-440-516-6	Sequence 6, Appli
32	64	100.0	160	4	US-10-440-516-7	Sequence 7, Appli
33	64	100.0	160	4	US-10-440-516-8	Sequence 8, Appli
34	64	100.0	160	4	US-10-440-516-11	Sequence 11, Appli
35	64	100.0	160	4	US-10-440-516-12	Sequence 12, Appli
36	64	100.0	160	4	US-10-440-516-15	Sequence 15, Appli
37	64	100.0	160	4	US-10-440-516-16	Sequence 16, Appli
38	64	100.0	160	4	US-10-440-516-19	Sequence 19, Appli
39	64	100.0	160	4	US-10-440-516-20	Sequence 20, Appli
40	64	100.0	160	4	US-10-440-516-23	Sequence 23, Appli
41	64	100.0	160	4	US-10-440-516-24	Sequence 24, Appli
42	64	100.0	160	4	US-10-440-516-25	Sequence 25, Appli
43	64	100.0	160	4	US-10-440-516-26	Sequence 26, Appli
44	64	100.0	160	4	US-10-440-516-27	Sequence 27, Appli
45	64	100.0	160	4	US-10-440-516-28	Sequence 28, Appli

ALIGNMENTS

RESULT 1
US-09-897-042-19
; Sequence 19, Application US/09897042
; Patent No. US20020018779A1
; GENERAL INFORMATION:
; APPLICANT: VALENTA, Rudolf
; APPLICANT: VRTALA, Susanne
; APPLICANT: VANGELISTA, Luca
; APPLICANT: EICHLER, Hans-Georg
; APPLICANT: SPERR, Wolfgang R.
; APPLICANT: VALENT, Peter
; APPLICANT: EBNER, Christof
; APPLICANT: KRAFT, Dietrich
; APPLICANT: GRONLUND, Hans
; TITLE OF INVENTION: NON-ANAPHYLACTIC FORMS OF ALLERGENS AND THEIR USE
; FILE REFERENCE: 1614-251P
; CURRENT APPLICATION NUMBER: US/09/897,042
; CURRENT FILING DATE: 2001-07-07
; NUMBER OF SEQ ID NOS: 21
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 19
; TYPE: PRT
; ORGANISM: Betula sp.
US-09-897-042-19

Query Match 100.0%; Score 64; DB 3; Length 12;
Best Local Similarity 100.0%; Pred. No. 0.00022;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
|||||
Db 1 GPGTIKKISFPE 12

RESULT 2
US-10-026-911-3
; Sequence 3, Application US/10026911
; Publication No. US20030078201A1
; GENERAL INFORMATION:
; APPLICANT: Focke, Margarete
; APPLICANT: Mahler, Vera
; APPLICANT: Sperr, Wolfgang R.
; APPLICANT: Valent, Peter
; APPLICANT: Kraft, Dietrich
; APPLICANT: Valenta, Rudolf
; TITLE OF INVENTION: Allergy Vaccines and Their Preparation
; FILE REFERENCE: 0273-0005
; CURRENT APPLICATION NUMBER: US/10/026,911
; CURRENT FILING DATE: 2002-07-24

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; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 31
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:
; OTHER INFORMATION: solvent-exposed peptide
US-10-026-911-3

Query Match          100.0%; Score 64; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.00059;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12
   |||||
Db 2 GPGTIKKISFPE 13

RESULT 3
US-10-799-514-5
; Sequence 5, Application US/10799514
; Publication No. US20040241178A1
; GENERAL INFORMATION:
; APPLICANT: Spertini, Francois
; APPLICANT: Cortesey, Blaise
; TITLE OF INVENTION: Allergen Peptide Fragments and Use Thereof
; FILE REFERENCE: 25720-502
; CURRENT APPLICATION NUMBER: US/10/799,514
; CURRENT FILING DATE: 2004-03-12
; PRIOR APPLICATION NUMBER: 60/455,004
; PRIOR FILING DATE: 2003-03-14
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 5
; LENGTH: 125
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: Peptide
US-10-799-514-5

Query Match          100.0%; Score 64; DB 5; Length 125;
Best Local Similarity 100.0%; Pred. No. 0.0026;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12
   |||||
Db 50 GPGTIKKISFPE 61

RESULT 4
US-09-981-009B-1
; Sequence 1, Application US/09981009B
; Publication No. US20030041354A1
; GENERAL INFORMATION:
; APPLICANT: Kjaerulff, Soren
; APPLICANT: Roggen, Erwin
; TITLE OF INVENTION: Transgenic Plants
; FILE REFERENCE: 10082.200-US
; CURRENT APPLICATION NUMBER: US/09/981,009B
; CURRENT FILING DATE: 2002-06-14
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula pendula
US-09-981-009B-1

Query Match          100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12
   |||||
Db 49 GPGTIKKISFPE 60

RESULT 5
US-09-847-208-34
; Sequence 34, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: IGE-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 34
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-34

Query Match          100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12
   |||||
Db 49 GPGTIKKISFPE 60

RESULT 6
US-09-847-208-37
; Sequence 37, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: IGE-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 37
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-37

Query Match          100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12
   |||||
Db 49 GPGTIKKISFPE 60

RESULT 7
US-09-847-208-38
; Sequence 38, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
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; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: IGE-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 38
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-38
Query Match      100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 GPGTIKKISFPE 12
        |||||
Db       49 GPGTIKKISFPE 60

RESULT 8
US-09-847-208-40
; Sequence 40, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: IGE-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 40
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-40
Query Match      100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 GPGTIKKISFPE 12
        |||||
Db       49 GPGTIKKISFPE 60

RESULT 9
US-09-957-806A-6
; Sequence 6, Application US/09957806A
; Publication No. US2005018146A1
; GENERAL INFORMATION:
; APPLICANT: Roggen, Erwin
; APPLICANT: Ernst, Steffen
; APPLICANT: Svendsen, Allan
; APPLICANT: Friis, Ebben
; APPLICANT: Osten, Claus
; TITLE OF INVENTION: PROTEIN VARIANTS HAVING MODIFIED IMMUNOGENICITY
; FILE REFERENCE: 10021.204-US
; CURRENT APPLICATION NUMBER: US/09/957,806A
; CURRENT FILING DATE: 2001-09-21
; NUMBER OF SEQ ID NOS: 248
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 6
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula pendula
US-09-957-806A-6
Query Match      100.0%; Score 64; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 GPGTIKKISFPE 12
        |||||
Db       49 GPGTIKKISFPE 60

RESULT 10
US-10-001-245-2
; Sequence 2, Application US/10001245
; Publication No. US20030175312A1
; GENERAL INFORMATION:
; APPLICANT: HOLM, Jens
; APPLICANT: IPSEN, Henrik
; APPLICANT: LARSEN, Jorgen N.
; APPLICANT: SPANGFORT, Michael D.
; TITLE OF INVENTION: No. US20030175312A1el mutant allergens
; FILE REFERENCE: 4305/IH942-US2
; CURRENT APPLICATION NUMBER: US/10/001,245
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/298,170
; PRIOR FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/249,361
; PRIOR FILING DATE: 2000-11-16
; NUMBER OF SEQ ID NOS: 217
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 2
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (5)..(5)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (42)..(42)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (45)..(45)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (78)..(78)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (103)..(103)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (123)..(123)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (134)..(134)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (156)..(156)
; OTHER INFORMATION:
; OTHER INFORMATION:
US-10-001-245-2
Query Match      100.0%; Score 64; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy      1 GPGTIKKISFPE 12
Db      49 GPGTIKKISFPE 60

RESULT 11
US-10-001-245-3
; Sequence 3, Application US/10001245
; Publication No. US20030175312A1
; GENERAL INFORMATION:
; APPLICANT: IPSEN, Henrik
; APPLICANT: LARSEN, Jorgen N.
; APPLICANT: SPANGFORT, Michael D.
; TITLE OF INVENTION: No. US20030175312A1el mutant allergens
; FILE REFERENCE: 4305/1H942-US2
; CURRENT APPLICATION NUMBER: US/10/001,245
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/298,170
; PRIOR FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/249,361
; PRIOR FILING DATE: 2000-11-16
; NUMBER OF SEQ ID NOS: 217
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa
; NAME/KEY: mutation
; LOCATION: (28)..(28)
; FEATURE:
; OTHER INFORMATION:
; NAME/KEY: mutation
; LOCATION: (45)..(45)
; FEATURE:
; OTHER INFORMATION:
; NAME/KEY: mutation
; LOCATION: (108)..(108)
; OTHER INFORMATION:
US-10-001-245-3
Query Match      100.0%; Score 64; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 GPGTIKKISFPE 12
Db      49 GPGTIKKISFPE 60

RESULT 12
US-10-001-245-4
; Sequence 4, Application US/10001245
; Publication No. US20030175312A1
; GENERAL INFORMATION:
; APPLICANT: HOLM, Jens
; APPLICANT: IPSEN, Henrik
; APPLICANT: LARSEN, Jorgen N.
; APPLICANT: SPANGFORT, Michael D.
; TITLE OF INVENTION: No. US20030175312A1el mutant allergens
; FILE REFERENCE: 4305/1H942-US2
; CURRENT APPLICATION NUMBER: US/10/001,245
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/298,170
; PRIOR FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/249,361
; PRIOR FILING DATE: 2000-11-16
```

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; NUMBER OF SEQ ID NOS: 217
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 4
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa
; NAME/KEY: mutation
; LOCATION: (5)..(5)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (45)..(45)
; OTHER INFORMATION:
; NAME/KEY: mutation
; LOCATION: (65)..(65)
; OTHER INFORMATION:
; NAME/KEY: mutation
; LOCATION: (97)..(97)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (134)..(134)
; OTHER INFORMATION:
US-10-001-245-4
Query Match      100.0%; Score 64; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 GPGTIKKISFPE 12
Db      49 GPGTIKKISFPE 60

RESULT 13
US-10-001-245-5
; Sequence 5, Application US/10001245
; Publication No. US20030175312A1
; GENERAL INFORMATION:
; APPLICANT: HOLM, Jens
; APPLICANT: IPSEN, Henrik
; APPLICANT: LARSEN, Jorgen N.
; APPLICANT: SPANGFORT, Michael D.
; TITLE OF INVENTION: No. US20030175312A1el mutant allergens
; FILE REFERENCE: 4305/1H942-US2
; CURRENT APPLICATION NUMBER: US/10/001,245
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/298,170
; PRIOR FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/249,361
; PRIOR FILING DATE: 2000-11-16
; NUMBER OF SEQ ID NOS: 217
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa
; NAME/KEY: mutation
; LOCATION: (16)..(16)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (28)..(28)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (32)..(32)
; OTHER INFORMATION:
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; NAME/KEY: mutation
; LOCATION: (103)..(103)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (108)..(108)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (152)..(152)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (153)..(153)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (155)..(155)
; OTHER INFORMATION:
; US-10-001-245-5
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Query Match 100.0%; Score 64; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 GPGTIKKISFPE 12
    |||||
Db 49 GPGTIKKISFPE 60
```

RESULT 14

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US-10-001-245-9
; Sequence 9, Application US/10001245
; Publication No. US20030175312A1
; GENERAL INFORMATION:
; APPLICANT: HOLM, Jens
; APPLICANT: IPSEN, Henrik
; APPLICANT: LARSEN, Jorgen N.
; APPLICANT: SPANGFORT, Michael D.
; TITLE OF INVENTION: No. US20030175312A1el mutant allergens
; FILE REFERENCE: 4305/1H942-US2
; CURRENT APPLICATION NUMBER: US/10/001,245
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/298,170
; PRIOR FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/249,361
; PRIOR FILING DATE: 2000-11-16
; NUMBER OF SEQ ID NOS: 217
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 9
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; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (28)..(28)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (32)..(32)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (65)..(65)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (96)..(96)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (97)..(97)
; OTHER INFORMATION:
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; FEATURE:
; NAME/KEY: mutation
; LOCATION: (108)..(108)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (109)..(109)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (125)..(125)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (127)..(127)
; OTHER INFORMATION:
; FEATURE:
; NAME/KEY: mutation
; LOCATION: (145)..(145)
; OTHER INFORMATION:
; US-10-001-245-9
```

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Query Match 100.0%; Score 64; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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```
QY 1 GPGTIKKISFPE 12
    |||||
Db 49 GPGTIKKISFPE 60
```

RESULT 15

```
US-10-001-245-92
; Sequence 92, Application US/10001245
; Publication No. US20030175312A1
; GENERAL INFORMATION:
; APPLICANT: HOLM, Jens
; APPLICANT: IPSEN, Henrik
; APPLICANT: LARSEN, Jorgen N.
; APPLICANT: SPANGFORT, Michael D.
; TITLE OF INVENTION: No. US20030175312A1el mutant allergens
; FILE REFERENCE: 4305/1H942-US2
; CURRENT APPLICATION NUMBER: US/10/001,245
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/298,170
; PRIOR FILING DATE: 2001-06-14
; PRIOR APPLICATION NUMBER: US 60/249,361
; PRIOR FILING DATE: 2000-11-16
; NUMBER OF SEQ ID NOS: 217
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 92
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa
; US-10-001-245-92
```

```
Query Match 100.0%; Score 64; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0034;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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```
QY 1 GPGTIKKISFPE 12
    |||||
Db 49 GPGTIKKISFPE 60
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OM protein - protein search, using sw model

Run on: January 28, 2006, 06:20:07 ; Search time 18.6667 Seconds
(without alignments)
53.149 Million cell updates/sec

Title: US-09-897-042-19
Perfect score: 64
Sequence: 1 GPGTIKISFPE 12

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Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
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2: /cgn2_6/prodata/1/iaa/6 COMB.pep.*
3: /cgn2_6/prodata/1/iaa/H COMB.pep.*
4: /cgn2_6/prodata/1/iaa/ECTUS COMB.pep.*
5: /cgn2_6/prodata/1/iaa/RE COMB.pep.*
6: /cgn2_6/prodata/1/iaa/backfiles1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	64	100.0	160	1	US-07-847-010-23
2	61	95.3	160	1	US-07-847-010-3
3	55	85.9	158	2	US-08-964-722-2
4	50	78.1	158	6	5312912-2
5	47	73.4	160	1	US-07-847-010-11
6	47	73.4	160	1	US-07-847-010-14
7	47	73.4	160	1	US-07-847-010-17
8	47	73.4	160	1	US-07-847-010-20
9	41	64.1	1198	2	US-09-252-991A-21016
10	40	62.5	1300	2	US-09-248-796A-27885
11	39	60.9	133	2	US-09-012-504A-18
12	39	60.9	581	2	US-09-107-532A-5385
13	39	60.9	592	2	US-09-830-230A-346
14	39	60.9	612	2	US-09-830-230A-345
15	38	59.4	405	2	US-09-134-000C-4680
16	38	59.4	471	1	US-08-399-889-24
17	38	59.4	471	2	US-09-167-364-24
18	38	59.4	471	2	US-09-439-897-2
19	38	59.4	3730	2	US-09-949-016-9908
20	37	57.8	60	2	US-09-540-236-2344
21	37	57.8	101	2	US-09-902-540-15306
22	37	57.8	300	2	US-09-982-616-9
23	37	57.8	680	2	US-09-489-039A-8422
24	37	57.8	763	2	US-09-583-110-3773
25	37	57.8	776	2	US-09-107-433-3635
26	37	57.8	793	1	US-08-468-558-5
27	37	57.8	793	2	US-08-676-444-5

28	37	57.8	1670	2	US-09-949-016-5883	Sequence 5883, Ap
29	36	56.2	74	2	US-09-248-796A-22915	Sequence 22915, A
30	36	56.2	163	2	US-09-861-451A-44	Sequence 44, Appl
31	36	56.2	168	2	US-09-605-703B-2820	Sequence 2820, Ap
32	36	56.2	447	2	US-09-109-204-3	Sequence 3, Appli
33	36	56.2	447	2	US-09-490-032-3	Sequence 3, Appli
34	36	56.2	453	2	US-09-252-991A-19829	Sequence 19829, A
35	36	56.2	608	2	US-09-464-377-2	Sequence 2, Appli
36	36	56.2	608	2	US-09-464-377-3	Sequence 3, Appli
37	35	54.7	84	2	US-09-248-796A-22781	Sequence 22781, A
38	35	54.7	131	2	US-09-641-638-654	Sequence 654, App
39	35	54.7	131	2	US-10-170-097-654	Sequence 654, App
40	35	54.7	133	2	US-09-012-515A-18	Sequence 18, Appl
41	35	54.7	133	2	US-08-360-144A-18	Sequence 18, Appl
42	35	54.7	133	2	US-09-012-399A-18	Sequence 18, Appl
43	35	54.7	133	4	PCT-US95-06722-18	Sequence 18, Appl
44	35	54.7	154	1	US-08-363-010-1	Sequence 1, Appli
45	35	54.7	154	1	US-08-911-434A-4	Sequence 4, Appli

ALIGNMENTS

RESULT 1
US-07-847-010-23
; Sequence 23, Application US/07847010
; Patent No. 5693495
; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikerstorfer, Arnold
; APPLICANT: Valentia, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
; APPLICANT: Rumpold, Helmut
; APPLICANT: Scheiner, Otto
; APPLICANT: Ebner, Christof
; APPLICANT: Ferreira, Fatima
; TITLE OF INVENTION: Allergens of Alder Pollen and
; TITLE OF INVENTION: Applications Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 23:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHEICAL: NO
; ORIGINAL SOURCE:

```
; ORGANISM: birch (Betula sp.)
; IMMEDIATE SOURCE:
; LIBRARY: POLLEN FROM ALLERCON AB, ENGELHOLM, SWEDEN
US-07-847-010-23

Query Match      100.0%; Score 64; DB 1; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.00051;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 GPGTIKKISFPE 12
      |||||||
Db      50 GPGTIKKISFPE 61

RESULT 2
US-07-847-010-3
; Sequence 3, Application US/07847010
; Patent No. 5693495
; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikerstorfer, Arnold
; APPLICANT: Valenta, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
; APPLICANT: Rumpold, Helmut
; APPLICANT: Scheiner, Otto
; APPLICANT: Ebner, Christof
; APPLICANT: Ferreira, Fatima
; TITLE OF INVENTION: Allergens of Alder Pollen and
; TITLE OF INVENTION: Applications Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C.
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; ORGANISM: Alder (Alnus sp.)
US-07-847-010-3

Query Match      95.3%; Score 61; DB 1; Length 160;
Best Local Similarity 91.7%; Pred. No. 0.0018;
Matches 11; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy      1 GPGTIKKISFPE 12
      |||||||
Db      50 GPGTIKKISFPE 61

; ORGANISM: birch (Betula sp.)
; IMMEDIATE SOURCE:
; LIBRARY: POLLEN FROM ALLERCON AB, ENGELHOLM, SWEDEN
US-07-847-010-23

Query Match      100.0%; Score 64; DB 1; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.00051;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 GPGTIKKISFPE 12
      |||||||
Db      50 GPGTIKKISFPE 61

RESULT 3
US-08-964-722-2
; Sequence 2, Application US/08964722A
; Patent No. 6184437
; GENERAL INFORMATION:
; APPLICANT: Sun, Samuel S.M.
; APPLICANT: Xiong, Liwen
; APPLICANT: Jing, Yuxiang
; APPLICANT: Liu, Bolin
; TITLE OF INVENTION: LYSINE RICH PROTEIN FROM WINGED BEAN
; FILE REFERENCE: 23461-20006.01
; CURRENT APPLICATION NUMBER: US/08/964,722A
; CURRENT FILING DATE: 1997-11-05
; EARLIER APPLICATION NUMBER: 60/002,918
; EARLIER FILING DATE: 1995-08-29
; EARLIER APPLICATION NUMBER: 60/004,266
; EARLIER FILING DATE: 1995-09-25
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 2
; TYPE: PRT
; LENGTH: 158
; ORGANISM: winged bean
US-08-964-722-2

Query Match      85.9%; Score 55; DB 2; Length 158;
Best Local Similarity 91.7%; Pred. No. 0.022;
Matches 11; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 GPGTIKKISFPE 12
      |||||||
Db      49 GPGTIKKISFVE 60

RESULT 4
5312912-2
; Patent No. 5312912
; APPLICANT: HADWIGER, LEE A.; CHIANG, CHIN C.; HOROVITZ, DANIEL A.
; TITLE OF INVENTION: PROCEDURES AND REGULATORY DNA SEQUENCES
; FOR GENETICALLY ENGINEERING DISEASE RESISTANCE AND OTHER
; INDUCIBLE TRAITS IN PLANTS
; NUMBER OF SEQUENCES: 9
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/393,301
; FILING DATE: 13-JUN-1989
; SEQ ID NO:2:
; LENGTH: 158
5312912-2

Query Match      78.1%; Score 50; DB 6; Length 158;
Best Local Similarity 75.0%; Pred. No. 0.18;
Matches 9; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy      1 GPGTIKKISFPE 12
      |||||||
Db      49 GPGTIKKLTFFE 60

RESULT 5
US-07-847-010-11
; Sequence 11, Application US/07847010
; Patent No. 5693495
; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikerstorfer, Arnold
; APPLICANT: Valenta, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
```

APPLICANT: Rumpold, Helmut
APPLICANT: Scheiner, Otto
APPLICANT: Ebner, Christof
APPLICANT: Ferreira, Fatima
TITLE OF INVENTION: Allergens of Alder Pollen and
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/847,010
FILING DATE: 01-JUN-1992
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jones III, Harry C
REGISTRATION NUMBER: 20,280
REFERENCE/DOCKET NUMBER: 6530-010
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 160 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: hazel (Corylus sp.)
IMMEDIATE SOURCE:
LIBRARY: POLLEN FROM ALLERGON AB, ENGELHOLM, SWEDEN
US-07-847-010-11

Query Match 73.4%; Score 47; DB 1; Length 160;
Best Local Similarity 75.0%; Pred. No. 0.66;
Matches 9; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
Db 50 GPGTIKNITFGE 61

RESULT 6
US-07-847-010-14
Sequence 14, Application US/07847010
Patent No. 5693495
GENERAL INFORMATION:
APPLICANT: Breiteneder, Heimo
APPLICANT: Reikerstorfer, Arnold
APPLICANT: Valenta, Rudolf
APPLICANT: Hoffmann - Sommergruber, Karin
APPLICANT: Breitenbach, Michael
APPLICANT: Rumpold, Helmut
APPLICANT: Scheiner, Otto
APPLICANT: Ebner, Christof
APPLICANT: Ferreira, Fatima
TITLE OF INVENTION: Allergens of Alder Pollen and
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds

STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/847,010
FILING DATE: 01-JUN-1992
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jones III, Harry C
REGISTRATION NUMBER: 20,280
REFERENCE/DOCKET NUMBER: 6530-010
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 160 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: hazel (Corylus sp.)
IMMEDIATE SOURCE:
LIBRARY: POLLEN FROM ALLERGON AB, ENGELHOLM, SWEDEN
US-07-847-010-14

Query Match 73.4%; Score 47; DB 1; Length 160;
Best Local Similarity 75.0%; Pred. No. 0.66;
Matches 9; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
Db 50 GPGTIKNITFGE 61

RESULT 7
US-07-847-010-17
Sequence 17, Application US/07847010
Patent No. 5693495
GENERAL INFORMATION:
APPLICANT: Breiteneder, Heimo
APPLICANT: Reikerstorfer, Arnold
APPLICANT: Valenta, Rudolf
APPLICANT: Hoffmann - Sommergruber, Karin
APPLICANT: Breitenbach, Michael
APPLICANT: Rumpold, Helmut
APPLICANT: Scheiner, Otto
APPLICANT: Ebner, Christof
APPLICANT: Ferreira, Fatima
TITLE OF INVENTION: Allergens of Alder Pollen and
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA: US/07/847,010
; APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; NAME: Jones III, Harry C
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; INFORMATION FOR SEQ ID NO: 17:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; ORGANISM: hazel (Corylus sp.)
; IMMEDIATE SOURCE:
; LIBRARY: POLLEN FROM ALLERCON AB, ENGELHOLM, SWEDEN
US-07-847-010-17

Query Match 73.4%; Score 47; DB 1; Length 160;
Best Local Similarity 75.0%; Pred. No. 0.66;
Matches 9; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12
||||| :||
Db 50 GPGTIKNITFGE 61

RESULT 8
US-07-847-010-20
; Sequence 20, Application US/07847010
; Patent No. 5693495
; GENERAL INFORMATION:
; APPLICANT: Breiteneder, Heimo
; APPLICANT: Reikerstorfer, Arnold
; APPLICANT: Valenta, Rudolf
; APPLICANT: Hoffmann - Sommergruber, Karin
; APPLICANT: Breitenbach, Michael
; APPLICANT: Kraft, Dietrich
; APPLICANT: Rumpold, Helmut
; APPLICANT: Scheiner, Otto
; APPLICANT: Ebner, Christof
; APPLICANT: Ferreira, Fatima
; TITLE OF INVENTION: Allergens of Alder Pollen and
; APPLICATION NUMBER: 23
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/847,010
; FILING DATE: 01-JUN-1992
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jones III, Harry C
; REGISTRATION NUMBER: 20,280
; REFERENCE/DOCKET NUMBER: 6530-010

; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 790-9090
; TELEFAX: (212) 869-9741/8864
; NAME: Pennie
; INFORMATION FOR SEQ ID NO: 20:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 160 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; ORGANISM: hazel (Corylus sp.)
; IMMEDIATE SOURCE:
; LIBRARY: POLLEN FROM ALLERCON AB, ENGELHOLM, SWEDEN
US-07-847-010-20

Query Match 73.4%; Score 47; DB 1; Length 160;
Best Local Similarity 75.0%; Pred. No. 0.66;
Matches 9; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 GPGTIKKISFPE 12
||||| :||
Db 50 GPGTIKNITFGE 61

RESULT 9
US-09-252-991A-21016
; Sequence 21016, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 21016
; LENGTH: 1198
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-21016

Query Match 64.1%; Score 41; DB 2; Length 1198;
Best Local Similarity 70.0%; Pred. No. 73;
Matches 7; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 GPGTIKKISF 10
||||| :||
Db 7 GPGTIRALSF 16

RESULT 10
US-09-248-796A-27885
; Sequence 27885, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
; FILE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208


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; SEQ ID NO 27885
; LENGTH: 190
; TYPE: PRT
; ORGANISM: Candida albicans
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: (2)
; OTHER INFORMATION: Identity of amino acid sequences at the above locations are unknown
US-09-248-796A-27885

Query Match 62.5%; Score 40; DB 2; Length 190;
Best Local Similarity 58.3%; Pred. No. 15;
Matches 7; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
DB 177 GPTVTEKINVP 188

RESULT 11
US-09-012-504A-18
; Sequence 18, Application US/09012504A
; Patent No. 6464974
; GENERAL INFORMATION:
; APPLICANT: Berlin, V.
; APPLICANT: Chiu, I.
; APPLICANT: Cottarel, G.
; APPLICANT: Damagnez, V.
; TITLE OF INVENTION: IMMUNOSUPPRESSANT TARGET PROTEINS
; FILE REFERENCE: APBI-P05-036
; CURRENT APPLICATION NUMBER: US/09/012,504A
; CURRENT FILING DATE: 1998-01-23
; PRIOR APPLICATION NUMBER: 08/360,144
; PRIOR FILING DATE: 1994-12-20
; PRIOR APPLICATION NUMBER: 08/250,795
; PRIOR FILING DATE: 1994-05-27
; NUMBER OF SEQ ID NOS: 35
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 18
; LENGTH: 133
; TYPE: PRT
; ORGANISM: Mammalian
US-09-012-504A-18

Query Match 60.9%; Score 39; DB 2; Length 133;
Best Local Similarity 58.3%; Pred. No. 16;
Matches 7; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
DB 46 GPTLTREISFQE 57

RESULT 12
US-09-107-532A-5385
; Sequence 5385, Application US/09107532A
; Patent No. 6583275
; GENERAL INFORMATION:
; APPLICANT: Lynn A Doucette-Stamm and David Bush
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
; NUMBER OF SEQUENCES: 7310
; CORRESPONDENCE ADDRESS:
; ADDRESS: GENOME THERAPEUTICS CORPORATION
; STREET: 100 Beaver Street
; CITY: Waltham
; STATE: Massachusetts
; COUNTRY: USA
; ZIP: 02354
; COMPUTER READABLE FORM:
; MEDIUM TYPE: CD/ROM ISO9660
; COMPUTER: PC
; OPERATING SYSTEM: <Unknown>

; SEQ ID NO 27885
; LENGTH: 190
; TYPE: PRT
; ORGANISM: Candida albicans
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: (2)
; OTHER INFORMATION: Identity of amino acid sequences at the above locations are unknown
US-09-248-796A-27885

Query Match 62.5%; Score 40; DB 2; Length 190;
Best Local Similarity 58.3%; Pred. No. 15;
Matches 7; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
DB 177 GPTVTEKINVP 188

RESULT 11
US-09-012-504A-18
; Sequence 18, Application US/09012504A
; Patent No. 6464974
; GENERAL INFORMATION:
; APPLICANT: Berlin, V.
; APPLICANT: Chiu, I.
; APPLICANT: Cottarel, G.
; APPLICANT: Damagnez, V.
; TITLE OF INVENTION: IMMUNOSUPPRESSANT TARGET PROTEINS
; FILE REFERENCE: APBI-P05-036
; CURRENT APPLICATION NUMBER: US/09/012,504A
; CURRENT FILING DATE: 1998-01-23
; PRIOR APPLICATION NUMBER: 08/360,144
; PRIOR FILING DATE: 1994-12-20
; PRIOR APPLICATION NUMBER: 08/250,795
; PRIOR FILING DATE: 1994-05-27
; NUMBER OF SEQ ID NOS: 35
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 18
; LENGTH: 133
; TYPE: PRT
; ORGANISM: Mammalian
US-09-012-504A-18

Query Match 60.9%; Score 39; DB 2; Length 133;
Best Local Similarity 58.3%; Pred. No. 16;
Matches 7; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
DB 46 GPTLTREISFQE 57

RESULT 12
US-09-107-532A-5385
; Sequence 5385, Application US/09107532A
; Patent No. 6583275
; GENERAL INFORMATION:
; APPLICANT: Lynn A Doucette-Stamm and David Bush
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
; NUMBER OF SEQUENCES: 7310
; CORRESPONDENCE ADDRESS:
; ADDRESS: GENOME THERAPEUTICS CORPORATION
; STREET: 100 Beaver Street
; CITY: Waltham
; STATE: Massachusetts
; COUNTRY: USA
; ZIP: 02354
; COMPUTER READABLE FORM:
; MEDIUM TYPE: CD/ROM ISO9660
; COMPUTER: PC
; OPERATING SYSTEM: <Unknown>

; SOFTWARE: ASCII
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/107,532A
; FILING DATE: 30-Jun-1998
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/085,598
; FILING DATE: 14 May 1998
; APPLICATION NUMBER: 60/051571
; FILING DATE: July 2, 1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Ariniello, Pamela Deneke
; REGISTRATION NUMBER: 40,489
; REFERENCE/DOCKET NUMBER: GTC-012
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (781)893-8277
; TELEFAX: (781)893-8277
; INFORMATION FOR SEQ ID NO: 5385:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 581 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; HYPOTHEICAL: YES
; ORIGINAL SOURCE:
; ORGANISM: Enterococcus faecium
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (B) LOCATION 1...581
; SEQUENCE DESCRIPTION: SEQ ID NO: 5385:
US-09-107-532A-5385

Query Match 60.9%; Score 39; DB 2; Length 581;
Best Local Similarity 58.3%; Pred. No. 78;
Matches 7; Conservative 1; Mismatches 4; Indels 0; Gaps 0;

QY 1 GPGTIKKISFPE 12
DB 352 GPGPAKKIRMPD 363

RESULT 13
US-09-830-230A-346
; Sequence 346, Application US/09830230A
; Patent No. 6902893
; GENERAL INFORMATION:
; APPLICANT: Human Genome Sciences, Inc.
; TITLE OF INVENTION: Lyme Disease Vaccines
; FILE REFERENCE: PB481US
; CURRENT APPLICATION NUMBER: US/09/830,230A
; CURRENT FILING DATE: 2001-09-27
; PRIOR APPLICATION NUMBER: PCT/US98/12718
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/057,483
; PRIOR FILING DATE: 1997-09-03
; PRIOR APPLICATION NUMBER: 60/053,344
; PRIOR FILING DATE: 1997-07-22
; PRIOR APPLICATION NUMBER: 60/053,377
; PRIOR FILING DATE: 1997-07-22
; PRIOR APPLICATION NUMBER: 60/050,359
; PRIOR FILING DATE: 1997-06-20
; NUMBER OF SEQ ID NOS: 756
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 346
; LENGTH: 592
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-830-230A-346

Query Match 60.9%; Score 39; DB 2; Length 592;
Best Local Similarity 88.9%; Pred. No. 79;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 3 GTIKKISFP 11
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Db 183 GFIKKISFP 191

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RESULT 14
US-09-830-230A-345
; Sequence 345, Application US/09830230A
; Patent No. 6902893
; GENERAL INFORMATION:
; APPLICANT: Human Genome Sciences, Inc.
; TITLE OF INVENTION: Lyme Disease Vaccines
; FILE REFERENCE: PB481US
; CURRENT APPLICATION NUMBER: US/09/830,230A
; CURRENT FILING DATE: 2001-09-27
; PRIOR APPLICATION NUMBER: PCT/US98/12718
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/057,483
; PRIOR FILING DATE: 1997-09-03
; PRIOR APPLICATION NUMBER: 60/053,344
; PRIOR FILING DATE: 1997-07-22
; PRIOR APPLICATION NUMBER: 60/053,377
; PRIOR FILING DATE: 1997-07-22
; PRIOR APPLICATION NUMBER: 60/050,359
; PRIOR FILING DATE: 1997-06-20
; NUMBER OF SEQ ID NOS: 756
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 345
; LENGTH: 612
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-830-230A-345

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Query Match      60.9%; Score 39; DB 2; Length 612;
Best Local Similarity 88.9%; Pred. No. 82;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
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Qy 3 GTIKISFP 11
Db 203 GFIIKISFP 211

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RESULT 15
US-09-134-000C-4680
; Sequence 4680, Application US/09134000C
; Patent No. 6617156
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
; TITLE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 032796-032
; CURRENT APPLICATION NUMBER: US/09/134,000C
; CURRENT FILING DATE: 1998-08-13
; PRIOR APPLICATION NUMBER: US 60/055,778
; PRIOR FILING DATE: 1997-08-15
; NUMBER OF SEQ ID NOS: 6812
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 4680
; LENGTH: 405
; TYPE: PRT
; ORGANISM: Enterococcus faecalis
US-09-134-000C-4680

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Query Match      59.4%; Score 38; DB 2; Length 405;
Best Local Similarity 54.5%; Pred. No. 80;
Matches 6: Conservative 4; Mismatches 1; Indels 0; Gaps 0;
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Qy      2 PGTIKKISFPE 12
        |||||:::|
Db     158 PGTIKRVSWAD 168
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Search completed: January 28, 2006, 06:24:30
Job time : 19.6667 secs

GenCore version 5.1.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.

OM protein - protein search, using sw model

Run on: January 28, 2006, 06:23:49 ; Search time 12.3333 Seconds
(without alignments)
10.536 Million cell updates/sec

Title: US-09-897-042-21
Perfect score: 68
Sequence: 1 KISFPEGFPFKY 12

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 75621 seqs, 10829074 residues

Total number of hits satisfying chosen parameters: 75621

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA.New.*
1: /cgn2_6/prodata/2/pubpaa/US08_NEW_PUB.pep.*
2: /cgn2_6/prodata/2/pubpaa/US06_NEW_PUB.pep.*
3: /cgn2_6/prodata/2/pubpaa/US07_NEW_PUB.pep.*
4: /cgn2_6/prodata/2/pubpaa/PCT_NEW_PUB.pep.*
5: /cgn2_6/prodata/2/pubpaa/US09_NEW_PUB.pep.*
6: /cgn2_6/prodata/2/pubpaa/US10_NEW_PUB.pep.*
7: /cgn2_6/prodata/2/pubpaa/US11_NEW_PUB.pep.*
8: /cgn2_6/prodata/2/pubpaa/US60_NEW_PUB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	68	100.0	172	7 US-11-102-883-18	Sequence 18, Appl
2	68	100.0	289	7 US-11-102-883-26	Sequence 26, Appl
3	68	100.0	300	7 US-11-102-883-6	Sequence 6, Appl
4	42	61.8	225	6 US-10-209-208-1	Sequence 1, Appl
5	42	61.8	225	6 US-10-209-208-4	Sequence 4, Appl
6	42	61.8	225	6 US-10-209-208-8	Sequence 8, Appl
7	42	61.8	225	6 US-10-209-208-20	Sequence 20, Appl
8	42	61.8	225	6 US-10-209-208-24	Sequence 24, Appl
9	42	61.8	225	6 US-10-209-208-79	Sequence 79, Appl
10	42	61.8	225	7 US-11-218-880-1	Sequence 1, Appl
11	42	61.8	225	7 US-11-218-880-4	Sequence 4, Appl
12	42	61.8	225	7 US-11-218-880-8	Sequence 8, Appl
13	42	61.8	225	7 US-11-218-880-20	Sequence 20, Appl
14	42	61.8	225	7 US-11-218-880-24	Sequence 24, Appl
15	42	61.8	225	7 US-11-100-988-2	Sequence 2, Appl
16	42	61.8	225	7 US-11-100-988-3	Sequence 3, Appl
17	42	61.8	225	7 US-11-100-988-5	Sequence 5, Appl
18	42	61.8	226	6 US-10-209-208-6	Sequence 6, Appl
19	42	61.8	226	7 US-11-218-880-6	Sequence 6, Appl
20	37	54.4	233	7 US-11-179-411-27	Sequence 27, Appl
21	37	54.4	233	7 US-11-179-411-33	Sequence 33, Appl
22	37	54.4	233	7 US-11-175-766-27	Sequence 27, Appl
23	37	54.4	233	7 US-11-175-766-33	Sequence 33, Appl
24	37	54.4	495	6 US-10-467-962B-93	Sequence 93, Appl
25	36	52.9	161	6 US-10-986-501-189	Sequence 189, Appl

26 36 52.9 238 7 US-11-052-554A-38 Sequence 38, Appl
27 35 51.5 343 6 US-10-131-826A-162 Sequence 162, Appl
28 35 51.5 607 7 US-11-024-959-381 Sequence 381, Appl
29 35 51.5 616 7 US-11-120-308-42 Sequence 42, Appl
30 35 51.5 1023 6 US-10-821-234-1377 Sequence 1377, Appl
31 35 51.5 1377 6 US-10-467-657-7922 Sequence 7922, Appl
32 34.5 50.7 216 7 US-11-082-389-32 Sequence 32, Appl
33 34 50.0 34 6 US-10-467-657-4010 Sequence 4010, Appl
34 34 50.0 108 6 US-10-821-234-1634 Sequence 1634, Appl
35 34 50.0 179 7 US-11-186-284-222 Sequence 222, Appl
36 34 50.0 199 6 US-10-821-234-1308 Sequence 1208, Appl
37 34 50.0 238 7 US-11-052-554A-51 Sequence 51, Appl
38 34 50.0 238 7 US-11-052-554A-52 Sequence 52, Appl
39 34 50.0 291 7 US-11-010-239-119 Sequence 119, Appl
40 34 50.0 300 7 US-11-151-601-17 Sequence 17, Appl
41 34 50.0 1185 6 US-10-877-346-7 Sequence 7, Appl
42 33 48.5 35 6 US-10-467-657-5572 Sequence 5572, Appl
43 33 48.5 49 7 US-11-145-861-307 Sequence 307, Appl
44 33 48.5 183 7 US-11-069-642-121 Sequence 121, Appl
45 33 48.5 833 6 US-10-467-657-3876 Sequence 3876, Appl

ALIGNMENTS

RESULT 1

US-11-102-883-18
; Sequence 18, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lampung, Norbert
; APPLICANT: Cramer, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods and
; TITLE OF INVENTION: Uses Thereof
; FILE REFERENCE: 03100234pa
; CURRENT APPLICATION NUMBER: US/11/102,883
; CURRENT FILING DATE: 2005-04-11
; PRIOR APPLICATION NUMBER: EP02022774.0
; PRIOR FILING DATE: 2002-10-11
; PRIOR APPLICATION NUMBER: PCT/EP2003/011190
; PRIOR FILING DATE: 2003-10-09
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 18
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Betula verrucosa
US-11-102-883-18

Query Match 100.0%; Score 68; DB 7; Length 172;
Best Local Similarity 100.0%; Pred. No. 6.3e-05;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12

Db 68 KISFPEGFPFKY 79

RESULT 2

US-11-102-883-26
; Sequence 26, Application US/11102883
; Publication No. US20050281816A1
; GENERAL INFORMATION:
; APPLICANT: Lampung, Norbert
; APPLICANT: Cramer, Reto
; APPLICANT: Fluckiger, Sabina
; APPLICANT: Daigle, Isabelle
; TITLE OF INVENTION: Modular Antigen Transporter Molecules (MAT Molecules) for
; TITLE OF INVENTION: Modulating Immune Reactions, Associated Constructs, Methods and
; TITLE OF INVENTION: Uses Thereof

; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; TITLE OF INVENTION: FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed polypeptide variant "mRFP1"
US-10-209-208-8

Query Match 61.8%; Score 42; DB 6; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
|:|||||:|:
Db 84 KLSFPEGFKWE 94

RESULT 7

US-10-209-208-20
; Sequence 20, Application US/10209208
; Publication No. US20050244921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; TITLE OF INVENTION: FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 20
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed with I125R
US-10-209-208-20

Query Match 61.8%; Score 42; DB 6; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
|:|||||:|:
Db 84 KLSFPEGFKWE 94

RESULT 8

US-10-209-208-24
; Sequence 24, Application US/10209208
; Publication No. US20050244921A1
; GENERAL INFORMATION:

; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; TITLE OF INVENTION: FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 24
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed polypeptide variant "T1" with I125R mutation
US-10-209-208-24

Query Match 61.8%; Score 42; DB 6; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
|:|||||:|:
Db 84 KLSFPEGFKWE 94

RESULT 9

US-10-209-208-79
; Sequence 79, Application US/10209208
; Publication No. US20050244921A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; APPLICANT: Geoffrey Baird
; TITLE OF INVENTION: FLUORESCENT PROTEIN VARIANTS AND METHODS
; TITLE OF INVENTION: FOR MAKING SAME
; FILE REFERENCE: UC083.1CP2CP2
; CURRENT APPLICATION NUMBER: US/10/209,208
; CURRENT FILING DATE: 2002-07-29
; PRIOR APPLICATION NUMBER: 10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; NUMBER OF SEQ ID NOS: 80
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 79
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed polypeptide variant "mRFP1.1"
US-10-209-208-79

Query Match 61.8%; Score 42; DB 6; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KISFPEGFPFK 11
|:|||||:|:
Db 84 KLSFPEGFKWE 94

RESULT 10

US-11-218-880-1

; Sequence 1, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Discosoma sp.
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(225)
; OTHER INFORMATION: wild-type DeRed
US-11-218-880-1

Query Match 61.8%; Score 42; DB 7; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFK 11
|:||||| :
Db 84 KLSFPEGFKWE 94

RESULT 11
US-11-218-880-4
; Sequence 4, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed polypeptide variant "T1"
US-11-218-880-4

Query Match 61.8%; Score 42; DB 7; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFK 11
|:||||| :
Db 84 KLSFPEGFKWE 94

RESULT 12
US-11-218-880-8
; Sequence 8, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed polypeptide variant "mRFP1"
US-11-218-880-8

Query Match 61.8%; Score 42; DB 7; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFK 11
|:||||| :
Db 84 KLSFPEGFKWE 94

RESULT 13
US-11-218-880-20
; Sequence 20, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Campbell, Robert
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; FILE REFERENCE: UC083.1CP2CP1
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/866,538
; PRIOR FILING DATE: 2001-05-24
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 20
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DeRed with I125R
US-11-218-880-20

Query Match 61.8%; Score 42; DB 7; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFK 11
|:||||| :
Db 84 KLSFPEGFKWE 94

Search completed: January 28, 2006, 06:44:48
Job time : 13.3333 secs

Sat Jan 28 18:36:01 2006

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RESULT 14
US-11-218-880-24
; Sequence 24, Application US/11218880
; Publication No. US20060003420A1
; GENERAL INFORMATION:
; APPLICANT: Tsien, Roger
; APPLICANT: Campbell, Robert
; TITLE OF INVENTION: MONOMERIC AND DIMERIC FLUORESCENT
; FILE REFERENCE: UC083.1CP2CPI
; CURRENT APPLICATION NUMBER: US/11/218,880
; CURRENT FILING DATE: 2005-09-01
; PRIOR FILING DATE: 2002-04-10
; PRIOR APPLICATION NUMBER: US/10/121,258
; PRIOR FILING DATE: 2001-02-26
; PRIOR APPLICATION NUMBER: 09/794,308
; PRIOR FILING DATE: 2001-05-24
; PRIOR APPLICATION NUMBER: 09/866,538
; NUMBER OF SEQ ID NOS: 78
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 24
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: DsRed polypeptide variant "T1" with I125R mutation
US-11-218-880-24

Query Match          61.8%; Score 42; DB 7; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 KISFPEGFPFK 11
       |:|||||:
Db      84 KLSFPEGFKWE 94

RESULT 15
US-11-100-988-2
; Sequence 2, Application US/11100988
; Publication No. US20060008878A1
; GENERAL INFORMATION:
; APPLICANT: Glick, Benjamin S.
; APPLICANT: Bevis, Brooke
; APPLICANT: Strongin, Daniel E.
; APPLICANT: Baker, David
; APPLICANT: Scalley-Kim, Michelle
; TITLE OF INVENTION: MONOMERIC RED FLUORESCENT PROTEINS
; FILE REFERENCE: 092234-9002-US01
; CURRENT APPLICATION NUMBER: US/11/100,988
; CURRENT FILING DATE: 2005-04-07
; PRIOR APPLICATION NUMBER: US 60/560,340
; PRIOR FILING DATE: 2004-04-07
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 225
; TYPE: PRT
; ORGANISM: Unknown
; FEATURE:
; OTHER INFORMATION: wild-type red fluorescent protein based on Discosoma species
US-11-100-988-2

Query Match          61.8%; Score 42; DB 7; Length 225;
Best Local Similarity 63.6%; Pred. No. 2;
Matches 7; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 KISFPEGFPFK 11
       |:|||||:
Db      84 KLSFPEGFKWE 94
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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: January 28, 2006, 06:23:08 ; Search time 79.6667 Seconds
(without alignments)
62.937 Million cell updates/sec

Title: US-09-897-042-21
Perfect score: 68
Sequence: 1 KISFPEGFPFKY 12

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA Main:
1: /cgn2_6/prodata/1/pubpaa/US07_PUBCOMB.pcp:*
2: /cgn2_6/prodata/1/pubpaa/US08_PUBCOMB.pcp:*
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5: /cgn2_6/prodata/1/pubpaa/US10B_PUBCOMB.pcp:*
6: /cgn2_6/prodata/1/pubpaa/US11_PUBCOMB.pcp:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	ID	Description
1	68	100.0	12	3	US-09-897-042-21
2	68	100.0	12	4	US-10-026-911-3
3	68	100.0	125	5	US-10-799-514-5
4	68	100.0	159	3	US-09-981-009B-1
5	68	100.0	159	3	US-09-847-208-34
6	68	100.0	159	3	US-09-847-208-38
7	68	100.0	159	3	US-09-847-208-40
8	68	100.0	159	3	US-09-957-806A-6
9	68	100.0	159	4	US-10-440-516-47
10	68	100.0	160	4	US-10-440-516-2
11	68	100.0	160	4	US-10-440-516-3
12	68	100.0	160	4	US-10-440-516-4
13	68	100.0	160	4	US-10-440-516-5
14	68	100.0	160	4	US-10-440-516-6
15	68	100.0	160	4	US-10-440-516-7
16	68	100.0	160	4	US-10-440-516-8
17	68	100.0	160	5	US-10-799-514-7
18	68	100.0	160	5	US-10-809-689-87
19	68	100.0	161	4	US-10-440-516-44
20	68	100.0	161	4	US-10-440-516-45
21	68	100.0	195	5	US-10-799-514-18
22	68	100.0	195	5	US-10-799-514-20
23	65	95.6	159	3	US-09-847-208-36
24	65	95.6	159	3	US-09-847-208-39
25	65	95.6	159	3	US-09-847-208-42
26	62	91.2	159	3	US-09-847-208-37
27	62	91.2	159	4	US-10-001-245-2

28	62	91.2	159	4	US-10-001-245-3	Sequence 3, Appli
29	62	91.2	159	4	US-10-001-245-5	Sequence 5, Appli
30	62	91.2	159	4	US-10-001-245-92	Sequence 92, Appli
31	62	91.2	159	4	US-10-719-553-37	Sequence 37, Appli
32	62	91.2	159	4	US-10-698-855-5	Sequence 5, Appli
33	62	91.2	160	4	US-10-001-245-1	Sequence 1, Appli
34	62	91.2	160	4	US-10-001-245-6	Sequence 6, Appli
35	62	91.2	160	4	US-10-001-245-8	Sequence 8, Appli
36	62	91.2	160	4	US-10-001-245-10	Sequence 10, Appli
37	62	91.2	160	4	US-10-440-516-1	Sequence 1, Appli
38	57	84.6	161	4	US-10-440-516-43	Sequence 43, Appli
39	57	83.8	159	3	US-09-847-208-8	Sequence 8, Appli
40	57	83.8	159	3	US-09-847-208-35	Sequence 35, Appli
41	57	83.8	159	3	US-09-847-208-41	Sequence 41, Appli
42	57	83.8	159	3	US-09-847-208-43	Sequence 43, Appli
43	57	83.8	159	4	US-10-001-245-4	Sequence 4, Appli
44	57	83.8	159	4	US-10-001-245-9	Sequence 9, Appli
45	57	83.8	160	4	US-10-001-245-7	Sequence 7, Appli

ALIGNMENTS

RESULT 1
US-09-897-042-21
; Sequence 21, Application US/09897042
; Patent No. US20020018779A1
; GENERAL INFORMATION:
; APPLICANT: VALENTA, Rudolf
; APPLICANT: VRTALIA, Susanne
; APPLICANT: VANGELISTA, Luca
; APPLICANT: EICHLER, Hans-Georg
; APPLICANT: SPERR, Wolfgang R.
; APPLICANT: VALENT, Peter
; APPLICANT: EBNER, Christof
; APPLICANT: KRAFT, Dietrich
; APPLICANT: GRONLUND, Hans
; TITLE OF INVENTION: NON-ANAPHYLACTIC FORMS OF ALLERGENS AND THEIR USE
; FILE REFERENCE: 1614-251P
; CURRENT APPLICATION NUMBER: US/09/897,042
; CURRENT FILING DATE: 2001-07-07
; NUMBER OF SEQ ID NOS: 21
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 21
; LENGTH: 12
; TYPE: PRT
; ORGANISM: Betula sp.
US-09-897-042-21

Query Match 100.0%; Score 68; DB 3; Length 12;
Best Local Similarity 100.0%; Pred. No. 0.00026;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12
|||||
Db 1 KISFPEGFPFKY 12

RESULT 2
US-10-026-911-3
; Sequence 3, Application US/10026911
; Publication No. US20030078201A1
; GENERAL INFORMATION:
; APPLICANT: Focke, Margarete
; APPLICANT: Mahler, Vera
; APPLICANT: Sperr, Wolfgang R.
; APPLICANT: Valent, Peter
; APPLICANT: Kraft, Dietrich
; APPLICANT: Valenta, Rudolf
; TITLE OF INVENTION: Allergy Vaccines and Their Preparation
; FILE REFERENCE: 0273-0005
; CURRENT APPLICATION NUMBER: US/10/026,911
; CURRENT FILING DATE: 2002-07-24

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; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 31
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:
; OTHER INFORMATION: solvent-exposed peptide
US-10-026-911-3

Query Match      100.0%; Score 68; DB 4; Length 31;
Best Local Similarity 100.0%; Pred. No. 0.00069;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12
Db 8 KISFPEGFPFKY 19

RESULT 3
US-10-799-514-5
; Sequence 5, Application US/10799514
; Publication No. US20040241178A1
; GENERAL INFORMATION:
; APPLICANT: Spertini, Francois
; APPLICANT: Cortesey, Blaise
; TITLE OF INVENTION: Allergen Peptide Fragments and Use Thereof
; FILE REFERENCE: 25720-502
; CURRENT APPLICATION NUMBER: US/10/799,514
; CURRENT FILING DATE: 2004-03-12
; PRIOR APPLICATION NUMBER: 60/455,004
; PRIOR FILING DATE: 2003-03-14
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 5
; LENGTH: 125
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: Peptide
US-10-799-514-5

Query Match      100.0%; Score 68; DB 5; Length 125;
Best Local Similarity 100.0%; Pred. No. 0.0028;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12
Db 56 KISFPEGFPFKY 67

RESULT 4
US-09-981-009B-1
; Sequence 1, Application US/09981009B
; Publication No. US20030041154A1
; GENERAL INFORMATION:
; APPLICANT: Kjaerulff, Soren
; APPLICANT: Roggen, Erwin
; TITLE OF INVENTION: Transgenic Plants
; FILE REFERENCE: 10082.200-US
; CURRENT APPLICATION NUMBER: US/09/981,009B
; CURRENT FILING DATE: 2002-06-14
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula pendula
US-09-981-009B-1

Query Match      100.0%; Score 68; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12
Db 55 KISFPEGFPFKY 66

RESULT 5
US-09-847-208-34
; Sequence 34, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: IGE-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 34
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-34

Query Match      100.0%; Score 68; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12
Db 55 KISFPEGFPFKY 66

RESULT 6
US-09-847-208-38
; Sequence 38, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: IGE-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 38
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (White birch) (Betula pendula)
US-09-847-208-38

Query Match      100.0%; Score 68; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12
Db 55 KISFPEGFPFKY 66

RESULT 7
US-09-847-208-40
; Sequence 40, Application US/09847208
; Publication No. US20030082190A1
; GENERAL INFORMATION:
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; APPLICANT: Saxon, Andrew
; APPLICANT: Zhang, Ke
; APPLICANT: Zhu, Daocheng
; TITLE OF INVENTION: FUSTON MOLECULES AND TREATMENT OF
; TITLE OF INVENTION: IGS-MEDIATED ALLERGIC DISEASES
; FILE REFERENCE: UC67.002A
; CURRENT APPLICATION NUMBER: US/09/847,208
; CURRENT FILING DATE: 2001-05-01
; NUMBER OF SEQ ID NOS: 177
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 40
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa (white birch) (Betula pendula)
; US-09-847-208-40

Query Match      100.0%; Score 68; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KISFPEGFPFKY 12
Db 55 KISFPEGFPFKY 66

RESULT 8
US-09-957-806A-6
; Sequence 6, Application US/09957806A
; Publication No. US20050181446A1
; GENERAL INFORMATION:
; APPLICANT: Roggen, Erwin
; APPLICANT: Ernst, Steffen
; APPLICANT: Svendsen, Allan
; APPLICANT: Friis, Esben
; APPLICANT: Osten, Claus
; TITLE OF INVENTION: PROTEIN VARIANTS HAVING MODIFIED IMMUNOGENICITY
; FILE REFERENCE: 10021.204-US
; CURRENT APPLICATION NUMBER: US/09/957.806A
; CURRENT FILING DATE: 2001-09-21
; NUMBER OF SEQ ID NOS: 248
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 6
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula pendula
; US-09-957-806A-6

Query Match      100.0%; Score 68; DB 3; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KISFPEGFPFKY 12
Db 55 KISFPEGFPFKY 66

RESULT 9
US-10-440-516-47
; Sequence 47, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 47
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen (amino acid deletion of residue 159)
; US-10-440-516-47

Query Match      100.0%; Score 68; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KISFPEGFPFKY 12
Db 55 KISFPEGFPFKY 66

RESULT 10
US-10-440-516-2
; Sequence 2, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; US-10-440-516-2

Query Match      100.0%; Score 68; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KISFPEGFPFKY 12
Db 56 KISFPEGFPFKY 67

RESULT 11
US-10-440-516-3
; Sequence 3, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 3
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
; US-10-440-516-3

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Query Match      100.0%; Score 68; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy 1 KISFPEGFFKY 12
Db 55 KISFPEGFFKY 66

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RESULT 12
US-10-440-516-4
; Sequence 4, Application US/10440516
; Publication No. US2004004338A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 4
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-4

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Query Match 100.0%; Score 68; DB 4; Length 160;
 Best Local Similarity 100.0%; Pred. No. 0.0036;
 Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	KISFPEGFPKY	12
Db	55	KISFPEGFPKY	66

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RESULT 13
US-10-440-516-5
; Sequence 5, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440.516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 5
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-5

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Query Match 100.0%; Score 68; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0

Qy 1 KISFPEGFFKY 12

Db 55 KISFEGFPFKY 66

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RESULT 14
US-10-440-516-6
; Sequence 6, Application US/10440516
; Publication No. US20040043438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 6
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-6

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Query Match      100.0%; Score 68; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12: Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy 1 KISFPEGFPKY 12
Db 55 KISFPEGFPKY 66

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RESULT 15
US-10-440-516-7
; Sequence 7, Application US/10440516
; Publication No. US200400438A1
; GENERAL INFORMATION:
; APPLICANT: Holm, Jens
; APPLICANT: Ferreras, Mercedes
; TITLE OF INVENTION: Allergen mutants
; FILE REFERENCE: 04305/100L446-US1
; CURRENT APPLICATION NUMBER: US/10/440,516
; CURRENT FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/381,440
; PRIOR FILING DATE: 2002-05-16
; NUMBER OF SEQ ID NOS: 89
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 7
; LENGTH: 160
; TYPE: PRP
; ORGANISM: Betula verrucosa
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (1)..(160)
; OTHER INFORMATION: Point mutated Bet v 1 allergen
US-10-440-516-7

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Query Match 100.0%; Score 68; DB 4; Length 160;
Best Local Similarity 100.0%; Pred. No. 0.0036;
Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KISFPEGFPFKY 12
Db 55 KISFPEGFPFKY 66

Search completed: January 28, 2006, 06:43:58

Job time : 79.6667 secs

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